

SEQUENCE LISTING

<110> Carozzi, Nadine
Hargiss, Tracy
Kozziel, Michael G.
Duck, Nicholas B.
Carr, Brian

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Methods for Its Use

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Ser	Asn	Ala	Ala	Cys	Val	Val	Tyr	Gly	Asn	Ser	Arg	Val	Asn	Val	Tyr	
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ata tca cct aac agt ggt tgt gat ttt acc ggt gat tac tat gag cga Ile Ser Pro Asn Ser Gly Cys Asp Phe Thr Gly Asp Tyr Tyr Glu Arg 225 230 235 240			720
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gat att atc gct ata ttt cca act tat gat ttt gag aaa tat cca ttg Asp Ile Ile Ala Ile Phe Pro Thr Tyr Asp Phe Glu Lys Tyr Pro Leu 290 295 300			912
cca aca cat gta gag ttg act agg gaa att tat aca gat gca gtg gga Pro Thr His Val Glu Leu Thr Arg Glu Ile Tyr Thr Asp Ala Val Gly 305 310 315 320			960
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Gly Trp Val Gly Thr Arg His Tyr Glu Asp Tyr Thr Lys Gly Asn Gly	
370 375 380	
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Ile Phe Gln Arg Met Ser Gly Thr Thr Ser Asn Asp Leu Arg Asn Ile	
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Asp Phe Gln Asn Ala Asp Val Tyr Lys Ile Thr Ser Leu Ala Ile Met	
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Asn Leu Val Gly Glu Thr Thr Ala Arg Pro Glu Tyr Arg Val Ser Lys	
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Ala Asp Phe Arg Arg Val Gly Gly Pro Asp Leu Asn Tyr Asp Ala Gly	
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Asn Asn Gly Leu Ser Arg Met Thr Ile Glu Ser Thr Phe Pro Leu Val	
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Leu His Ser Asn Gly Val Arg Gly Pro Ser His Arg Leu Ser Asn Ala	
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Ala Cys Val Val Tyr Gly Asn Ser Arg Val Asn Val Tyr Gly Trp Thr	
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cat aca agt tta aaa cgt gaa aat ata att gaa gcc aat caa att aca	1536
His Thr Ser Leu Lys Arg Glu Asn Ile Ile Glu Ala Asn Gln Ile Thr	
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caa ata ccg gcg gtg aag agt tat tac ctt caa aat tat ctt gct aat	1584
Gln Ile Pro Ala Val Lys Ser Tyr Tyr Leu Gln Asn Tyr Leu Ala Asn	
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Ala Tyr Thr Tyr Val Ile Lys Gly Thr His Thr Gly Gly Asp Leu Ile	
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Gly Ile Arg Leu Ile Ile Asn Asn Lys Thr Ala Gly Gln Ser Tyr Arg	
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Ile Arg Phe Arg Tyr Ala Ala Asp Lys Ala Ala Phe Phe Ser Val Tyr	
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Leu Tyr Pro Gly Gly Trp Gly Ser Asn Arg Phe Val Ser Leu Glu Lys	
595 600 605	
tct tac tct gga aat tat gac gat tta aaa tat agt gat ttt aaa ttc	1872
Ser Tyr Ser Gly Asn Tyr Asp Asp Leu Lys Tyr Ser Asp Phe Lys Phe	
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Ala Glu Ile Ile Thr Pro Pro Leu Pro Ser Ser Asn Ile Gln Met Asp	
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Val Glu Met Gln Ala Asn Ser Phe Gln Ser Asp Val Asn Val Val Leu	
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Asp Lys Ile Glu Phe Leu Pro Ser Asn Thr Thr Thr Leu Glu Tyr Glu	
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Glu Ile Trp Glu Ser Val Glu Thr Ile Thr Ser Ile Gly Ile Asn Leu	
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Ile Glu Phe Val Ile Glu Pro Ser Leu Gly Gly Ile Asn Thr Leu Leu	
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Leu Ser Ile Cys Asp Leu Leu Ser Ile Ile Arg Lys Glu Val Ala Asp	
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Ser Val Leu Ser Asp Ala Ile Ala Asp Phe Asp Gly Lys Leu Lys Asn	
115 120 125	
Tyr Arg Glu Tyr Tyr Leu Ser Tyr Leu Gly Ala Trp Leu Lys Asp Gly	
130 135 140	

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Tyr	Phe	Lys	Leu	Ser	Glu	Arg	Asp	Phe	Asn	Glu	Ile	Leu	Gly	Gly	Ser
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Leu	Ser	Arg	Asn	Asn	Ala	Gln	Val	Leu	Leu	Leu	Pro	Thr	Phe	Ala	Gln
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Ala	Ala	Asn	Val	Gln	Leu	Leu	Leu	Leu	Arg	Asp	Ala	Val	Gln	Tyr	Lys
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Ile	Ser	Pro	Asn	Ser	Gly	Cys	Asp	Phe	Thr	Gly	Asp	Tyr	Tyr	Glu	Arg
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Leu	Lys	Cys	Lys	Thr	Ala	Glu	Tyr	Thr	Asn	Tyr	Cys	Leu	Tyr	Trp	Tyr
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Gln	Val	Gly	Leu	Asn	Gln	Ile	Lys	Gln	Gly	Gly	Thr	Gly	Ala	Asp	Thr
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Trp	Ser	Lys	Phe	Asn	Lys	Phe	Arg	Arg	Glu	Met	Thr	Leu	Ala	Val	Leu
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Asp	Ile	Ile	Ala	Ile	Phe	Pro	Thr	Tyr	Asp	Phe	Glu	Lys	Tyr	Pro	Leu
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Pro	Thr	His	Val	Glu	Leu	Thr	Arg	Glu	Ile	Tyr	Thr	Asp	Ala	Val	Gly
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Tyr	Ser	Ser	Gly	Thr	Tyr	Ser	Trp	Leu	Arg	Asn	Trp	Pro	Asn	Thr	Phe
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Asn	Gly	Leu	Glu	Ala	Asn	Gly	Thr	Arg	Gly	Pro	Gly	Leu	Val	Thr	Trp
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Ile	Phe	Gln	Arg	Met	Ser	Gly	Thr	Thr	Ser	Asn	Asp	Leu	Arg	Asn	Ile
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Asp	Phe	Gln	Asn	Ala	Asp	Val	Tyr	Lys	Ile	Thr	Ser	Leu	Ala	Ile	Met
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Ala	Asp	Phe	Arg	Arg	Val	Gly	Gly	Pro	Asp	Leu	Asn	Tyr	Asp	Ala	Gly
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Leu	His	Ser	Asn	Gly	Val	Arg	Gly	Pro	Ser	His	Arg	Leu	Ser	Asn	Ala
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Ala	Cys	Val	Val	Tyr	Gly	Asn	Ser	Arg	Val	Asn	Val	Tyr	Gly	Trp	Thr
				485						490					495
His	Thr	Ser	Leu	Lys	Arg	Glu	Asn	Ile	Ile	Glu	Ala	Asn	Gln	Ile	Thr
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Gln	Ile	Pro	Ala	Val	Lys	Ser	Tyr	Tyr	Leu	Gln	Asn	Tyr	Leu	Ala	Asn
		515						520					525		
Ala	Tyr	Thr	Tyr	Val	Ile	Lys	Gly	Thr	His	Thr	Gly	Gly	Asp	Leu	Ile
	530					535					540				
Arg	Phe	Leu	Arg	Thr	Lys	Ser	Glu	Tyr	Asn	Ala	Val	Tyr	Ala	Gly	Gly
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Gly	Ile	Arg	Leu	Ile	Ile	Asn	Asn	Lys	Thr	Ala	Gly	Gln	Ser	Tyr	Arg
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Ile	Arg	Phe	Arg	Tyr	Ala	Ala	Asp	Lys	Ala	Ala	Phe	Phe	Ser	Val	Tyr
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Leu	Tyr	Pro	Gly	Gly	Trp	Gly	Ser	Asn	Arg	Phe	Val	Ser	Leu	Glu	Lys

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	610					615					620				
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	625				630					635				640	
Val	Glu	Met	Gln	Ala	Asn	Ser	Phe	Gln	Ser	Asp	Val	Asn	Val	Val	Leu
			645						650				655		
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Thr Thr Asp Tyr Glu Ile Asp Gln Ala Ala Ile Ser Ile Glu Cys Met	
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Ser Asp Glu Gln Asn Pro Gln Glu Lys Ile Met Leu Trp Asp Glu Ile	
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aaa ctg gca aaa caa ctt agt cag tct cgt aat cta ctc caa aat gga	192
Lys Leu Ala Lys Gln Leu Ser Gln Ser Arg Asn Leu Leu Gln Asn Gly	
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gac ttt tct ggg aat gat tgg aca ttc ggt aat gat att atc ata gga	240
Asp Phe Ser Gly Asn Asp Trp Thr Phe Gly Asn Asp Ile Ile Ile Gly	
65 70 75 80	
tcc aat aat cct att ttt aaa gga aaa ttt cta cag atg cgt gga gca	288
Ser Asn Asn Pro Ile Phe Lys Gly Lys Phe Leu Gln Met Arg Gly Ala	
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cga gac ata tat gga act cta ttt cca acc tat atc tgt caa aaa ata	336
Arg Asp Ile Tyr Gly Thr Leu Phe Pro Thr Tyr Ile Cys Gln Lys Ile	
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gat gag tct aaa tta aaa cca tat aca cgt tat cga gta aga ggg ttt	384
Asp Glu Ser Lys Leu Lys Pro Tyr Thr Arg Tyr Arg Val Arg Gly Phe	
115 120 125	

gtg gga agt agt aaa gat ttg aaa tta atg gta aca cgt tac ggg aaa	432
Val Gly Ser Ser Lys Asp Leu Lys Leu Met Val Thr Arg Tyr Gly Lys	
130 135 140	
gaa att gat gct atc atg aat gtt cca aat gat ttg gcc tat atg cag	480
Glu Ile Asp Ala Ile Met Asn Val Pro Asn Asp Leu Ala Tyr Met Gln	
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cct aat cct tca tgt gga gat tat cgc tgt gaa tca tcg tct cag tat	528
Pro Asn Pro Ser Cys Gly Asp Tyr Arg Cys Glu Ser Ser Ser Gln Tyr	
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gtg agc caa ggg tat cct aca cca aca gat gga tat gct ccc gat atg	576
Val Ser Gln Gly Tyr Pro Thr Pro Thr Asp Gly Tyr Ala Pro Asp Met	
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Tyr Ala Cys Pro Gln Asn Ile Asp Arg Lys His Val Lys Cys His Asp	
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Arg His Pro Phe Asp Phe His Ile Asp Thr Gly Glu Val Asp Thr Asn	
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Tyr Ala Thr Val Gly Asn Leu Glu Val Ile Glu Glu Gly Pro Leu Thr	
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His Met Glu Lys Lys Arg Trp Glu Thr Gln Gln Ala Tyr Asp Pro Ala	
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Lys Gln Ala Val Asp Ala Leu Phe Thr Asn Glu Gln Glu Leu His Tyr	
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His Ile Thr Leu Asp His Ile Gln Asn Ala Asp Arg Leu Val Gln Ser	
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Ile Pro Tyr Val Tyr His Asn Trp Leu Pro Asn Ala Pro Gly Met Asn	
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Asn Leu His Ala Gln Asp His His Gly Tyr Val Leu Arg Val Ile Ala	
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Gly Lys Gln Glu Thr Leu Lys Phe Thr Ser Cys Glu Glu Gly Tyr Met	
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Thr Lys Thr Val Glu Val Phe Pro Glu Ser Asp Arg Val Arg Ile Glu	
450 455 460	
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Ile Gly Glu Thr Glu Gly Thr Phe Tyr Ile Asp Ser Ile Glu Leu Leu	
465 470 475 480	
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Cys Met Gln Gly Tyr Asp Asn Asn Asn Asn Leu His Thr Gly Asn Met	
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Tyr Tyr Gln Gly Tyr Thr Asn Asn Tyr Asn Gln Asp Ser Ser Asn Met	
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Tyr Asn Gln Asn Tyr Thr Asn Asn Asp Asp Leu His Ser Gly Cys Thr	
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 <213> Bacillus thuringiensis

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Val	Ser	Gln	Gly	Tyr	Pro	Thr	Pro	Thr	Asp	Gly	Tyr	Ala	Pro	Asp	Met
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Tyr	Ala	Cys	Pro	Gln	Asn	Ile	Asp	Arg	Lys	His	Val	Lys	Cys	His	Asp
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Tyr	Ala	Thr	Val	Gly	Asn	Leu	Glu	Val	Ile	Glu	Glu	Gly	Pro	Leu	Thr
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Gly	Glu	Ala	Leu	Ala	His	Val	Lys	Gln	Lys	Glu	Lys	Lys	Trp	Lys	Gln
		260					265						270		
His	Met	Glu	Lys	Lys	Arg	Trp	Glu	Thr	Gln	Gln	Ala	Tyr	Asp	Pro	Ala
	275						280					285			
Lys	Gln	Ala	Val	Asp	Ala	Leu	Phe	Thr	Asn	Glu	Gln	Glu	Leu	His	Tyr
	290					295					300				
His	Ile	Thr	Leu	Asp	His	Ile	Gln	Asn	Ala	Asp	Arg	Leu	Val	Gln	Ser
305					310					315					320
Ile	Pro	Tyr	Val	Tyr	His	Asn	Trp	Leu	Pro	Asn	Ala	Pro	Gly	Met	Asn
			325						330					335	
Tyr	Asp	Val	Tyr	Gln	Glu	Leu	Asn	Ala	Arg	Ile	Met	Gln	Gly	Tyr	Asn
		340						345					350		
Leu	Tyr	Asp	Ala	Arg	Asn	Val	Ile	Thr	Asn	Gly	Asp	Phe	Thr	Gln	Gly
	355					360						365			
Leu	Gln	Gly	Trp	His	Ala	Thr	Gly	Asn	Ala	Ala	Val	Gln	Gln	Met	Asp
	370					375					380				
Gly	Ala	Ser	Val	Leu	Val	Leu	Ser	Asn	Trp	Ser	Ala	Gly	Val	Ser	Gln
385					390					395					400
Asn	Leu	His	Ala	Gln	Asp	His	His	Gly	Tyr	Val	Leu	Arg	Val	Ile	Ala
				405					410					415	

Lys	Lys	Glu	Gly	Pro	Gly	Lys	Gly	Tyr	Val	Thr	Met	Met	Asp	Cys	Asn
			420					425					430		
Gly	Lys	Gln	Glu	Thr	Leu	Lys	Phe	Thr	Ser	Cys	Glu	Glu	Gly	Tyr	Met
		435					440					445			
Thr	Lys	Thr	Val	Glu	Val	Phe	Pro	Glu	Ser	Asp	Arg	Val	Arg	Ile	Glu
	450					455					460				
Ile	Gly	Glu	Thr	Glu	Gly	Thr	Phe	Tyr	Ile	Asp	Ser	Ile	Glu	Leu	Leu
465					470					475					480
Cys	Met	Gln	Gly	Tyr	Asp	Asn	Asn	Asn	Asn	Leu	His	Thr	Gly	Asn	Met
				485					490					495	
Tyr	Glu	Gln	Ser	Tyr	Asn	Gly	Asn	Tyr	Asn	Gln	Asn	Thr	Ser	Asp	Val
		500						505					510		
Tyr	Tyr	Gln	Gly	Tyr	Thr	Asn	Asn	Tyr	Asn	Gln	Asp	Ser	Ser	Asn	Met
	515						520					525			
Tyr	Asn	Gln	Asn	Tyr	Thr	Asn	Asn	Asp	Asp	Leu	His	Ser	Gly	Cys	Thr
	530					535					540				
Cys	Asn	Gln	Gly	His	Asn	Ser	Gly	Cys	Thr	Cys	Asn	Gln	Gly	Tyr	Asn
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Arg															

<210> 8

<211> 1176

<212> PRT

<213> *Bacillus thuringiensis*

<400> 8

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		20						25				30			
Tyr	Thr	Pro	Ile	Asp	Ile	Ser	Leu	Ser	Leu	Thr	Gln	Phe	Leu	Leu	Ser
	35					40					45				
Glu	Phe	Val	Pro	Gly	Ala	Gly	Phe	Val	Leu	Gly	Leu	Val	Asp	Ile	Ile
	50					55				60					
Trp	Gly	Ile	Phe	Gly	Pro	Ser	Gln	Trp	Asp	Ala	Phe	Pro	Val	Gln	Ile
65				70					75					80	
Glu	Gln	Leu	Ile	Asn	Gln	Arg	Ile	Glu	Glu	Phe	Ala	Arg	Asn	Gln	Ala
				85				90					95		
Ile	Ser	Arg	Leu	Glu	Gly	Leu	Ser	Asn	Leu	Tyr	Gln	Ile	Tyr	Ala	Glu
		100				105					110				
Ser	Phe	Arg	Glu	Trp	Glu	Ala	Asp	Pro	Thr	Asn	Pro	Ala	Leu	Arg	Glu
	115					120					125				
Glu	Met	Arg	Ile	Gln	Phe	Asn	Asp	Met	Asn	Ser	Ala	Leu	Thr	Thr	Ala
	130					135				140					
Ile	Pro	Leu	Leu	Ala	Val	Gln	Asn	Tyr	Gln	Val	Pro	Leu	Leu	Ser	Val
145				150					155					160	
Tyr	Val	Gln	Ala	Ala	Asn	Leu	His	Leu	Ser	Val	Leu	Arg	Asp	Val	Ser
			165					170					175		
Val	Phe	Gly	Gln	Arg	Trp	Gly	Phe	Asp	Ala	Ala	Thr	Ile	Asn	Ser	Arg
	180					185						190			
Tyr	Asn	Asp	Leu	Thr	Arg	Leu	Ile	Gly	Asn	Tyr	Thr	Asp	Tyr	Ala	Val
	195					200					205				
Arg	Trp	Tyr	Asn	Thr	Gly	Leu	Glu	Arg	Val	Trp	Gly	Pro	Asp	Ser	Arg
	210				215					220					
Asp	Trp	Val	Arg	Tyr	Asn	Gln	Phe	Arg	Arg	Glu	Leu	Thr	Leu	Thr	Val

225					230					235					240
Leu	Asp	Ile	Val	Ala	Leu	Phe	Ser	Asn	Tyr	Asp	Ser	Arg	Arg	Tyr	Pro
				245					250					255	
Ile	Arg	Thr	Val	Ser	Gln	Leu	Thr	Arg	Glu	Ile	Tyr	Thr	Asn	Pro	Val
			260					265					270		
Leu	Glu	Asn	Phe	Asp	Gly	Ser	Phe	Arg	Gly	Met	Ala	Gln	Arg	Ile	Glu
		275					280					285			
Gln	Asn	Ile	Arg	Gln	Pro	His	Leu	Met	Asp	Ile	Leu	Asn	Ser	Ile	Thr
	290					295				300					
Ile	Tyr	Thr	Asp	Val	His	Arg	Gly	Phe	Asn	Tyr	Trp	Ser	Gly	His	Gln
305					310					315				320	
Ile	Thr	Ala	Ser	Pro	Val	Gly	Phe	Ser	Gly	Pro	Glu	Phe	Ala	Phe	Pro
			325						330					335	
Leu	Phe	Gly	Asn	Ala	Gly	Asn	Ala	Ala	Pro	Pro	Val	Leu	Val	Ser	Leu
		340					345					350			
Thr	Gly	Leu	Gly	Ile	Phe	Arg	Thr	Leu	Ser	Ser	Pro	Leu	Tyr	Arg	Arg
	355					360						365			
Ile	Ile	Leu	Gly	Ser	Gly	Pro	Asn	Asn	Gln	Glu	Leu	Phe	Val	Leu	Asp
	370				375					380					
Gly	Thr	Glu	Phe	Ser	Phe	Ala	Ser	Leu	Thr	Thr	Asn	Leu	Pro	Ser	Thr
385					390					395				400	
Ile	Tyr	Arg	Gln	Arg	Gly	Thr	Val	Asp	Ser	Leu	Asp	Val	Ile	Pro	Pro
			405					410					415		
Gln	Asp	Asn	Ser	Val	Pro	Pro	Arg	Ala	Gly	Phe	Ser	His	Arg	Leu	Ser
		420					425					430			
His	Val	Thr	Met	Leu	Ser	Gln	Ala	Ala	Gly	Ala	Val	Tyr	Thr	Leu	Arg
	435					440				445					
Ala	Pro	Thr	Phe	Ser	Trp	Gln	His	Arg	Ser	Ala	Glu	Phe	Asn	Asn	Ile
	450				455					460					
Ile	Pro	Ser	Ser	Gln	Ile	Thr	Gln	Ile	Pro	Leu	Thr	Lys	Ser	Thr	Asn
465					470					475				480	
Leu	Gly	Ser	Gly	Thr	Ser	Val	Val	Lys	Gly	Pro	Gly	Phe	Thr	Gly	Gly
			485					490					495		
Asp	Ile	Leu	Arg	Arg	Thr	Ser	Pro	Gly	Gln	Ile	Ser	Thr	Leu	Arg	Val
		500					505					510			
Asn	Ile	Thr	Ala	Pro	Leu	Ser	Gln	Arg	Tyr	Arg	Val	Arg	Ile	Arg	Tyr
	515					520					525				
Ala	Ser	Thr	Thr	Asn	Leu	Gln	Phe	His	Thr	Ser	Ile	Asp	Gly	Arg	Pro
	530				535					540					
Ile	Asn	Gln	Gly	Asn	Phe	Ser	Ala	Thr	Met	Ser	Ser	Gly	Ser	Asn	Leu
545				550					555				560		
Gln	Ser	Gly	Ser	Phe	Arg	Thr	Val	Gly	Phe	Thr	Thr	Pro	Phe	Asn	Phe
			565					570					575		
Ser	Asn	Gly	Ser	Ser	Val	Phe	Thr	Leu	Ser	Ala	His	Val	Phe	Asn	Ser
		580					585					590			
Gly	Asn	Glu	Val	Tyr	Ile	Asp	Arg	Ile	Glu	Phe	Val	Pro	Ala	Glu	Val
	595					600					605				
Thr	Phe	Glu	Ala	Glu	Tyr	Asp	Leu	Glu	Arg	Ala	Gln	Lys	Ala	Val	Asn
	610				615					620					
Glu	Leu	Phe	Thr	Ser	Ser	Asn	Gln	Ile	Gly	Leu	Lys	Thr	Asp	Val	Thr
625					630				635					640	
Asp	Tyr	His	Ile	Asp	Gln	Val	Ser	Asn	Leu	Val	Glu	Cys	Leu	Ser	Asp
			645					650					655		
Glu	Phe	Cys	Leu	Asp	Glu	Lys	Gln	Glu	Leu	Ser	Glu	Lys	Val	Lys	His
		660					665					670			
Ala	Lys	Arg	Leu	Ser	Asp	Glu	Arg	Asn	Leu	Leu	Gln	Asp	Pro	Asn	Phe
	675					680						685			

Arg	Gly	Ile	Asn	Arg	Gln	Leu	Asp	Arg	Gly	Trp	Arg	Gly	Ser	Thr	Asp	
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Ile	Thr	Ile	Gln	Gly	Gly	Asp	Asp	Val	Phe	Lys	Glu	Asn	Tyr	Val	Thr	
705					710					715					720	
Leu	Leu	Gly	Thr	Phe	Asp	Glu	Cys	Tyr	Pro	Thr	Tyr	Leu	Tyr	Gln	Lys	
				725					730						735	
Ile	Asp	Glu	Ser	Lys	Leu	Lys	Ala	Tyr	Thr	Arg	Tyr	Gln	Leu	Arg	Gly	
			740					745					750			
Tyr	Ile	Glu	Asp	Ser	Gln	Asp	Leu	Glu	Ile	Tyr	Leu	Ile	Arg	Tyr	Asn	
	755						760					765				
Ala	Lys	His	Glu	Thr	Val	Asn	Val	Pro	Gly	Thr	Gly	Ser	Leu	Trp	Pro	
	770					775					780					
Leu	Ser	Ala	Gln	Ser	Pro	Ile	Gly	Lys	Cys	Gly	Glu	Pro	Asn	Arg	Cys	
785					790					795					800	
Ala	Pro	His	Leu	Glu	Trp	Asn	Pro	Asp	Leu	Asp	Cys	Ser	Cys	Arg	Asp	
				805					810						815	
Gly	Glu	Lys	Cys	Ala	His	His	Ser	His	His	Phe	Ser	Leu	Asp	Ile	Asp	
			820					825						830		
Val	Gly	Cys	Thr	Asp	Leu	Asn	Glu	Asp	Leu	Gly	Val	Trp	Val	Ile	Phe	
	835						840					845				
Lys	Ile	Lys	Thr	Gln	Asp	Gly	His	Ala	Arg	Leu	Gly	Asn	Leu	Glu	Phe	
	850					855					860					
Leu	Glu	Glu	Lys	Pro	Leu	Val	Gly	Glu	Ala	Leu	Ala	Arg	Val	Lys	Arg	
865					870					875					880	
Ala	Glu	Lys	Lys	Trp	Arg	Asp	Lys	Arg	Glu	Lys	Leu	Glu	Trp	Glu	Thr	
				885					890						895	
Asn	Ile	Val	Tyr	Lys	Glu	Ala	Lys	Glu	Ser	Val	Asp	Ala	Leu	Phe	Val	
		900						905					910			
Asn	Ser	Gln	Tyr	Asp	Gln	Leu	Gln	Ala	Asp	Thr	Asn	Ile	Ala	Met	Ile	
	915						920						925			
His	Ala	Ala	Asp	Lys	Arg	Val	His	Ser	Ile	Arg	Glu	Ala	Tyr	Leu	Pro	
	930					935						940				
Glu	Leu	Ser	Val	Ile	Pro	Gly	Val	Asn	Ala	Ala	Ile	Phe	Glu	Glu	Leu	
945					950					955					960	
Glu	Gly	Arg	Ile	Phe	Thr	Ala	Phe	Ser	Leu	Tyr	Asp	Ala	Arg	Asn	Val	
				965					970						975	
Ile	Lys	Asn	Gly	Asp	Phe	Asn	Asn	Gly	Leu	Ser	Cys	Trp	Asn	Val	Lys	
			980					985						990		
Gly	His	Val	Asp	Val	Glu	Glu	Gln	Asn	Asn	Gln	Arg	Ser	Val	Leu	Val	
	995						1000					1005				
Val	Pro	Glu	Trp	Glu	Ala	Glu	Val	Ser	Gln	Glu	Val	Arg	Val	Cys	Pro	
	1010					1015					1020					
Gly	Arg	Gly	Tyr	Ile	Leu	Arg	Val	Thr	Ala	Tyr	Lys	Glu	Gly	Tyr	Gly	
1025					1030					1035					1040	
Glu	Gly	Cys	Val	Thr	Ile	His	Glu	Ile	Glu	Asn	Asn	Thr	Asp	Glu	Leu	
				1045					1050					1055		
Lys	Phe	Ser	Asn	Cys	Val	Glu	Glu	Glu	Ile	Tyr	Pro	Asn	Asn	Thr	Val	
		1060						1065					1070			
Thr	Cys	Asn	Asp	Tyr	Thr	Val	Asn	Gln	Glu	Glu	Tyr	Gly	Gly	Ala	Tyr	
	1075						1080					1085				
Thr	Ser	Arg	Asn	Arg	Gly	Tyr	Asn	Glu	Ala	Pro	Ser	Val	Pro	Ala	Asp	
	1090					1095					1100					
Tyr	Ala	Ser	Val	Tyr	Glu	Glu	Lys	Ser	Tyr	Thr	Asp	Gly	Arg	Arg	Glu	
1105					1110					1115					1120	
Asn	Pro	Cys	Glu	Phe	Asn	Arg	Gly	Tyr	Arg	Asp	Tyr	Thr	Pro	Leu	Pro	
				1125					1130					1135		
Val	Gly	Tyr	Val	Thr	Lys	Glu	Leu	Glu	Tyr	Phe	Pro	Glu	Thr	Asp	Lys	

	1140		1145		1150										
Val	Trp	Ile	Glu	Ile	Gly	Glu	Thr	Glu	Gly	Thr	Phe	Ile	Val	Asp	Ser
	1155		1160		1165										
Val	Glu	Leu	Leu	Leu	Met	Glu	Glu								
	1170					1175									

<210> 9
 <211> 1178
 <212> PRT
 <213> *Bacillus thuringiensis*

<400> 9

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			20					25					30		
Tyr	Thr	Pro	Ile	Asp	Ile	Ser	Leu	Ser	Leu	Thr	Gln	Phe	Leu	Leu	Ser
		35				40						45			
Glu	Phe	Val	Pro	Gly	Ala	Gly	Phe	Val	Leu	Gly	Leu	Val	Asp	Ile	Ile
	50					55					60				
Trp	Gly	Ile	Phe	Gly	Pro	Ser	Gln	Trp	Asp	Ala	Phe	Leu	Val	Gln	Ile
65					70				75					80	
Glu	Gln	Leu	Ile	Asn	Gln	Arg	Ile	Glu	Glu	Phe	Ala	Arg	Asn	Gln	Ala
				85					90					95	
Ile	Ser	Arg	Leu	Glu	Gly	Leu	Ser	Asn	Leu	Tyr	Gln	Ile	Tyr	Ala	Glu
			100					105					110		
Ser	Phe	Arg	Glu	Trp	Glu	Ala	Asp	Pro	Thr	Asn	Pro	Ala	Leu	Arg	Glu
		115					120					125			
Glu	Met	Arg	Ile	Gln	Phe	Asn	Asp	Met	Asn	Ser	Ala	Leu	Thr	Thr	Ala
	130					135					140				
Ile	Pro	Leu	Phe	Ala	Val	Gln	Asn	Tyr	Gln	Val	Pro	Leu	Leu	Ser	Val
145					150					155				160	
Tyr	Val	Gln	Ala	Ala	Asn	Leu	His	Leu	Ser	Val	Leu	Arg	Asp	Val	Ser
				165					170					175	
Val	Phe	Gly	Gln	Arg	Trp	Gly	Phe	Asp	Ala	Ala	Thr	Ile	Asn	Ser	Arg
			180					185					190		
Tyr	Asn	Asp	Leu	Thr	Arg	Leu	Ile	Gly	Asn	Tyr	Thr	Asp	Tyr	Ala	Val
		195					200					205			
Arg	Trp	Tyr	Asn	Thr	Gly	Leu	Glu	Arg	Val	Trp	Gly	Pro	Asp	Ser	Arg
	210					215					220				
Asp	Trp	Val	Arg	Tyr	Asn	Gln	Phe	Arg	Arg	Glu	Leu	Thr	Leu	Thr	Val
225					230					235				240	
Leu	Asp	Ile	Val	Ala	Leu	Phe	Pro	Asn	Tyr	Asp	Ser	Arg	Arg	Tyr	Pro
				245					250					255	
Ile	Arg	Thr	Val	Ser	Gln	Leu	Thr	Arg	Glu	Ile	Tyr	Thr	Asn	Pro	Val
			260					265					270		
Leu	Glu	Asn	Phe	Asp	Gly	Ser	Phe	Arg	Gly	Ser	Ala	Gln	Gly	Ile	Glu
		275					280					285			
Arg	Ser	Ile	Arg	Ser	Pro	His	Leu	Met	Asp	Ile	Leu	Asn	Ser	Ile	Thr
	290					295					300				
Ile	Tyr	Thr	Asp	Ala	His	Arg	Gly	Tyr	Tyr	Tyr	Trp	Ser	Gly	His	Gln
305					310					315				320	
Ile	Met	Ala	Ser	Pro	Val	Gly	Phe	Ser	Gly	Pro	Glu	Phe	Thr	Phe	Pro
				325					330					335	
Leu	Tyr	Gly	Thr	Met	Gly	Asn	Ala	Ala	Pro	Gln	Gln	Arg	Ile	Val	Ala
			340					345					350		

Gln	Leu	Gly	Gln	Gly	Val	Tyr	Arg	Thr	Leu	Ser	Ser	Thr	Leu	Tyr	Arg
		355					360					365			
Arg	Pro	Phe	Asn	Ile	Gly	Ile	Asn	Asn	Gln	Gln	Leu	Ser	Val	Leu	Asp
		370				375					380				
Gly	Thr	Glu	Phe	Ala	Tyr	Gly	Thr	Ser	Ser	Asn	Leu	Pro	Ser	Ala	Val
385					390					395					400
Tyr	Arg	Lys	Ser	Gly	Thr	Val	Asp	Ser	Leu	Asp	Glu	Ile	Pro	Pro	Gln
			405						410					415	
Asn	Asn	Asn	Val	Pro	Pro	Arg	Gln	Gly	Phe	Ser	His	Arg	Leu	Ser	His
			420					425					430		
Val	Ser	Met	Phe	Arg	Ser	Gly	Phe	Ser	Asn	Ser	Ser	Val	Ser	Ile	Ile
		435				440						445			
Arg	Ala	Pro	Met	Phe	Ser	Trp	Ile	His	Arg	Ser	Ala	Glu	Phe	Asn	Asn
		450				455					460				
Ile	Ile	Ala	Ser	Asp	Ser	Ile	Thr	Gln	Ile	Pro	Ala	Val	Lys	Gly	Asn
465					470					475					480
Phe	Leu	Phe	Asn	Gly	Ser	Val	Ile	Ser	Gly	Pro	Gly	Phe	Thr	Gly	Gly
			485						490					495	
Asp	Leu	Val	Arg	Leu	Asn	Ser	Ser	Gly	Asn	Asn	Ile	Gln	Asn	Arg	Gly
			500					505				510			
Tyr	Ile	Glu	Val	Pro	Ile	His	Phe	Pro	Ser	Thr	Ser	Thr	Arg	Tyr	Arg
		515					520					525			
Val	Arg	Val	Arg	Tyr	Ala	Ser	Val	Thr	Pro	Ile	His	Leu	Asn	Val	Asn
		530				535					540				
Trp	Gly	Asn	Ser	Ser	Ile	Phe	Ser	Asn	Thr	Val	Pro	Ala	Thr	Ala	Thr
545					550					555					560
Ser	Leu	Asp	Asn	Leu	Gln	Ser	Ser	Asp	Phe	Gly	Tyr	Phe	Glu	Ser	Ala
			565					570						575	
Asn	Ala	Phe	Thr	Ser	Ser	Leu	Gly	Asn	Ile	Val	Gly	Val	Arg	Asn	Phe
			580					585					590		
Ser	Gly	Thr	Ala	Gly	Val	Ile	Ile	Asp	Arg	Phe	Glu	Phe	Ile	Pro	Val
		595					600					605			
Thr	Ala	Thr	Leu	Glu	Ala	Glu	Tyr	Asn	Leu	Glu	Arg	Ala	Gln	Lys	Ala
		610				615					620				
Val	Asn	Ala	Leu	Phe	Thr	Ser	Thr	Asn	Gln	Leu	Gly	Leu	Lys	Thr	Asn
625					630					635					640
Val	Thr	Asp	Tyr	His	Ile	Asp	Gln	Val	Ser	Asn	Leu	Val	Thr	Tyr	Leu
			645					650						655	
Ser	Asp	Glu	Phe	Cys	Leu	Asp	Glu	Lys	Arg	Glu	Leu	Ser	Glu	Lys	Val
			660				665						670		
Lys	His	Ala	Lys	Arg	Leu	Ser	Asp	Glu	Arg	Asn	Leu	Leu	Gln	Asp	Ser
		675					680					685			
Asn	Phe	Lys	Asp	Ile	Asn	Arg	Gln	Pro	Glu	Arg	Gly	Trp	Gly	Gly	Ser
		690				695					700				
Thr	Gly	Ile	Thr	Ile	Gln	Gly	Gly	Asp	Asp	Val	Phe	Lys	Glu	Asn	Tyr
705					710					715					720
Val	Thr	Leu	Ser	Gly	Thr	Phe	Asp	Glu	Cys	Tyr	Pro	Thr	Tyr	Leu	Tyr
			725					730						735	
Gln	Lys	Ile	Asp	Glu	Ser	Lys	Leu	Lys	Ala	Phe	Thr	Arg	Tyr	Gln	Leu
			740					745					750		
Arg	Gly	Tyr	Ile	Glu	Asp	Ser	Gln	Asp	Leu	Glu	Ile	Tyr	Leu	Ile	Arg
		755					760					765			
Tyr	Asn	Ala	Lys	His	Glu	Thr	Val	Asn	Val	Pro	Gly	Thr	Gly	Ser	Leu
		770				775					780				
Trp	Pro	Leu	Ser	Ala	Gln	Ser	Pro	Ile	Gly	Lys	Cys	Gly	Glu	Pro	Asn
785					790					795					800
Arg	Cys	Ala	Pro	His	Leu	Glu	Trp	Asn	Pro	Asp	Leu	Asp	Cys	Ser	Cys

				805					810					815			
Arg	Asp	Gly	Glu	Lys	Cys	Ala	His	His	Ser	His	His	Phe	Ser	Leu	Asp		
			820						825					830			
Ile	Asp	Val	Gly	Cys	Thr	Asp	Leu	Asn	Glu	Asp	Leu	Gly	Val	Trp	Val		
		835						840					845				
Ile	Phe	Lys	Ile	Lys	Thr	Gln	Asp	Gly	His	Ala	Arg	Leu	Gly	Asn	Leu		
	850					855					860						
Glu	Phe	Leu	Glu	Glu	Lys	Pro	Leu	Val	Gly	Glu	Ala	Leu	Ala	Arg	Val		
865					870					875					880		
Lys	Arg	Ala	Glu	Lys	Trp	Arg	Asp	Lys	Arg	Glu	Lys	Leu	Glu	Trp			
			885						890					895			
Glu	Thr	Asn	Ile	Val	Tyr	Lys	Glu	Ala	Lys	Glu	Ser	Val	Asp	Ala	Leu		
		900						905					910				
Phe	Val	Asn	Ser	Gln	Tyr	Asp	Gln	Leu	Gln	Ala	Asp	Thr	Asn	Ile	Ala		
	915						920					925					
Met	Ile	His	Ala	Ala	Asp	Lys	Arg	Val	His	Ser	Ile	Arg	Glu	Ala	Tyr		
	930					935					940						
Leu	Pro	Glu	Leu	Ser	Val	Ile	Pro	Gly	Val	Asn	Ala	Ala	Ile	Phe	Glu		
945					950					955					960		
Glu	Leu	Glu	Gly	Arg	Ile	Phe	Thr	Ala	Phe	Ser	Leu	Tyr	Asp	Ala	Arg		
			965					970						975			
Asn	Val	Ile	Lys	Asn	Gly	Asp	Phe	Asn	Asn	Gly	Leu	Ser	Cys	Trp	Asn		
		980					985						990				
Val	Lys	Gly	His	Val	Asp	Val	Glu	Glu	Gln	Asn	Asn	Gln	Arg	Ser	Val		
	995						1000					1005					
Leu	Val	Val	Pro	Glu	Trp	Glu	Ala	Glu	Val	Ser	Gln	Glu	Val	Arg	Val		
	1010					1015					1020						
Cys	Pro	Gly	Arg	Gly	Tyr	Ile	Leu	Arg	Val	Thr	Ala	Tyr	Lys	Glu	Gly		
1025					1030					1035				1040			
Tyr	Gly	Glu	Gly	Cys	Val	Thr	Ile	His	Glu	Ile	Glu	Asn	Asn	Thr	Asp		
			1045						1050					1055			
Glu	Leu	Lys	Phe	Ser	Asn	Cys	Val	Glu	Glu	Glu	Ile	Tyr	Pro	Asn	Asn		
		1060						1065				1070					
Thr	Val	Thr	Cys	Asn	Asp	Tyr	Thr	Val	Asn	Gln	Glu	Glu	Tyr	Gly	Gly		
	1075						1080					1085					
Ala	Tyr	Thr	Ser	Arg	Asn	Arg	Gly	Tyr	Asn	Glu	Ala	Pro	Ser	Val	Pro		
	1090					1095					1100						
Ala	Asp	Tyr	Ala	Ser	Val	Tyr	Glu	Glu	Lys	Ser	Tyr	Thr	Asp	Gly	Arg		
1105					1110					1115				1120			
Arg	Glu	Asn	Pro	Cys	Glu	Phe	Asn	Arg	Gly	Tyr	Arg	Asp	Tyr	Thr	Pro		
			1125						1130					1135			
Leu	Pro	Val	Gly	Tyr	Val	Thr	Lys	Glu	Leu	Glu	Tyr	Phe	Pro	Glu	Thr		
		1140						1145				1150					
Asp	Lys	Val	Trp	Ile	Glu	Ile	Gly	Glu	Thr	Glu	Gly	Thr	Phe	Ile	Val		
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Asp	Ser	Val	Glu	Leu	Leu	Leu	Met	Glu	Glu								
	1170						1175										

<210> 10

<211> 719

<212> PRT

<213> Bacillus thuringiensis

<400> 10

Met	Lys	Leu	Lys	Asn	Gln	Asp	Lys	His	Gln	Ser	Phe	Ser	Ser	Asn	Ala
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Lys	Val	Asp	Lys	Ile	Ser	Thr	Asp	Ser	Leu	Lys	Asn	Glu	Thr	Asp	Ile	
			20					25					30			
Glu	Leu	Gln	Asn	Ile	Asn	His	Glu	Asp	Cys	Leu	Lys	Met	Ser	Glu	Tyr	
		35					40					45				
Glu	Asn	Val	Glu	Pro	Phe	Val	Ser	Ala	Ser	Thr	Ile	Gln	Thr	Gly	Ile	
		50				55					60					
Gly	Ile	Ala	Gly	Lys	Ile	Leu	Gly	Thr	Leu	Gly	Val	Pro	Phe	Ala	Gly	
65					70					75					80	
Gln	Val	Ala	Ser	Leu	Tyr	Ser	Phe	Ile	Leu	Gly	Glu	Leu	Trp	Pro	Lys	
			85						90					95		
Gly	Lys	Asn	Gln	Trp	Glu	Ile	Phe	Met	Glu	His	Val	Glu	Glu	Ile	Ile	
			100					105					110			
Asn	Gln	Lys	Ile	Ser	Thr	Tyr	Ala	Arg	Asn	Lys	Ala	Leu	Thr	Asp	Leu	
		115					120					125				
Lys	Gly	Leu	Gly	Asp	Ala	Leu	Ala	Val	Tyr	His	Asp	Ser	Leu	Glu	Ser	
		130				135					140					
Trp	Val	Gly	Asn	Arg	Asn	Asn	Thr	Arg	Ala	Arg	Ser	Val	Val	Lys	Ser	
145					150					155					160	
Gln	Tyr	Ile	Ala	Leu	Glu	Leu	Met	Phe	Val	Gln	Lys	Leu	Pro	Ser	Phe	
			165					170						175		
Ala	Val	Ser	Gly	Glu	Glu	Val	Pro	Leu	Leu	Pro	Ile	Tyr	Ala	Gln	Ala	
			180					185					190			
Ala	Asn	Leu	His	Leu	Leu	Leu	Leu	Arg	Asp	Ala	Ser	Ile	Phe	Gly	Lys	
		195					200					205				
Glu	Trp	Gly	Leu	Ser	Ser	Ser	Glu	Ile	Ser	Thr	Phe	Tyr	Asn	Arg	Gln	
		210				215					220					
Val	Glu	Arg	Ala	Gly	Asp	Tyr	Ser	Asp	His	Cys	Val	Lys	Trp	Tyr	Ser	
225					230					235					240	
Thr	Gly	Leu	Asn	Asn	Leu	Arg	Gly	Thr	Asn	Ala	Glu	Ser	Trp	Val	Arg	
			245						250					255		
Tyr	Asn	Gln	Phe	Arg	Arg	Asp	Met	Thr	Leu	Met	Val	Leu	Asp	Leu	Val	
		260						265					270			
Ala	Leu	Phe	Pro	Ser	Tyr	Asp	Thr	Gln	Met	Tyr	Pro	Ile	Lys	Thr	Thr	
		275					280					285				
Ala	Gln	Leu	Thr	Arg	Glu	Val	Tyr	Thr	Asp	Ala	Ile	Gly	Thr	Val	His	
		290				295					300					
Pro	His	Pro	Ser	Phe	Thr	Ser	Thr	Thr	Trp	Tyr	Asn	Asn	Asn	Ala	Pro	
305					310					315					320	
Ser	Phe	Ser	Ala	Ile	Glu	Ala	Ala	Val	Val	Arg	Asn	Pro	His	Leu	Leu	
			325					330						335		
Asp	Phe	Leu	Glu	Gln	Val	Thr	Ile	Tyr	Ser	Leu	Leu	Ser	Arg	Trp	Ser	
		340						345					350			
Asn	Thr	Gln	Tyr	Met	Asn	Met	Trp	Gly	Gly	His	Lys	Leu	Glu	Phe	Arg	
		355					360					365				
Thr	Ile	Gly	Gly	Thr	Leu	Asn	Ile	Ser	Thr	Gln	Gly	Ser	Thr	Asn	Thr	
		370				375					380					
Ser	Ile	Asn	Pro	Val	Thr	Leu	Pro	Phe	Thr	Ser	Arg	Asp	Val	Tyr	Arg	
385					390					395					400	
Thr	Glu	Ser	Leu	Ala	Gly	Leu	Asn	Leu	Phe	Leu	Thr	Gln	Pro	Val	Asn	
			405						410					415		
Gly	Val	Pro	Arg	Val	Asp	Phe	His	Trp	Lys	Phe	Val	Thr	His	Pro	Ile	
		420						425					430			
Ala	Ser	Asp	Asn	Phe	Tyr	Tyr	Pro	Gly	Tyr	Ala	Gly	Ile	Gly	Thr	Gln	
		435					440					445				
Leu	Gln	Asp	Ser	Glu	Asn	Glu	Leu	Pro	Pro	Glu	Ala	Thr	Gly	Gln	Pro	
		450				455					460					
Asn	Tyr	Glu	Ser	Tyr	Ser	His	Arg	Leu	Ser	His	Ile	Gly	Leu	Ile	Ser	

465					470					475				480	
Ala	Ser	His	Val	Lys	Ala	Leu	Val	Tyr	Ser	Trp	Thr	His	Arg	Ser	Ala
				485					490					495	
Asp	Arg	Thr	Asn	Thr	Ile	Glu	Pro	Asn	Ser	Ile	Thr	Gln	Ile	Pro	Leu
			500					505					510		
Val	Lys	Ala	Phe	Asn	Leu	Ser	Ser	Gly	Ala	Ala	Val	Val	Arg	Gly	Pro
		515					520					525			
Gly	Phe	Thr	Gly	Gly	Asp	Ile	Leu	Arg	Arg	Thr	Asn	Thr	Gly	Thr	Phe
	530					535					540				
Gly	Asp	Ile	Arg	Val	Asn	Ile	Asn	Pro	Pro	Phe	Ala	Gln	Arg	Tyr	Arg
545					550					555					560
Val	Arg	Ile	Arg	Tyr	Ala	Ser	Thr	Thr	Asp	Leu	Gln	Phe	His	Thr	Ser
				565					570					575	
Ile	Asn	Gly	Lys	Ala	Ile	Asn	Gln	Gly	Asn	Phe	Ser	Ala	Thr	Met	Asn
			580					585					590		
Arg	Gly	Glu	Asp	Leu	Asp	Tyr	Lys	Thr	Phe	Arg	Thr	Val	Gly	Phe	Thr
		595					600					605			
Thr	Pro	Phe	Ser	Phe	Leu	Asp	Val	Gln	Ser	Thr	Phe	Thr	Ile	Gly	Ala
	610					615					620				
Trp	Asn	Phe	Ser	Ser	Gly	Asn	Glu	Val	Tyr	Ile	Asp	Arg	Ile	Glu	Phe
625					630					635					640
Val	Pro	Val	Glu	Val	Thr	Tyr	Glu	Ala	Glu	Tyr	Asp	Phe	Glu	Lys	Ala
				645					650					655	
Gln	Glu	Lys	Val	Thr	Ala	Leu	Phe	Thr	Ser	Thr	Asn	Pro	Arg	Gly	Leu
			660					665					670		
Lys	Thr	Asp	Val	Lys	Asp	Tyr	His	Ile	Asp	Gln	Val	Ser	Asn	Leu	Val
		675					680					685			
Glu	Ser	Leu	Ser	Asp	Glu	Phe	Tyr	Leu	Asp	Glu	Lys	Arg	Glu	Leu	Phe
	690					695					700				
Glu	Ile	Val	Lys	Tyr	Ala	Lys	Gln	Leu	His	Ile	Glu	Arg	Asn	Met	
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 <212> PRT
 <213> Bacillus thuringiensis

<400> 11															
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Asn	Val	Val	Ala	His	Asp	Pro	Phe	Ser	Phe	Glu	His	Lys	Ser	Leu	Asp
			20					25					30		
Thr	Ile	Gln	Lys	Glu	Trp	Met	Glu	Trp	Lys	Arg	Thr	Asp	His	Ser	Leu
	35						40					45			
Tyr	Val	Ala	Pro	Val	Val	Gly	Thr	Val	Ser	Ser	Phe	Leu	Leu	Lys	Lys
	50					55					60				
Val	Gly	Ser	Leu	Ile	Gly	Lys	Arg	Ile	Leu	Ser	Glu	Leu	Trp	Gly	Ile
65					70					75					80
Ile	Phe	Pro	Ser	Gly	Ser	Thr	Asn	Leu	Met	Gln	Asp	Ile	Leu	Arg	Glu
				85					90					95	
Thr	Glu	Gln	Phe	Leu	Asn	Gln	Arg	Leu	Asn	Thr	Asp	Thr	Leu	Ala	Arg
			100					105					110		
Val	Asn	Ala	Glu	Leu	Ile	Gly	Leu	Gln	Ala	Asn	Ile	Arg	Glu	Phe	Asn
		115					120					125			
Gln	Gln	Val	Asp	Asn	Phe	Leu	Asn	Pro	Thr	Gln	Asn	Pro	Val	Pro	Leu
	130					135					140				

Ser	Ile	Thr	Ser	Ser	Val	Asn	Thr	Met	Gln	Gln	Leu	Phe	Leu	Asn	Arg
145					150				155					160	
Leu	Pro	Gln	Phe	Gln	Ile	Gln	Gly	Tyr	Gln	Leu	Leu	Leu	Leu	Pro	Leu
				165					170					175	
Phe	Ala	Gln	Ala	Ala	Asn	Met	His	Leu	Ser	Phe	Ile	Arg	Asp	Val	Ile
			180					185					190		
Leu	Asn	Ala	Asp	Glu	Trp	Gly	Ile	Ser	Ala	Ala	Thr	Leu	Arg	Thr	Tyr
		195					200					205			
Arg	Asp	Tyr	Leu	Arg	Asn	Tyr	Thr	Arg	Asp	Tyr	Ser	Asn	Tyr	Cys	Ile
	210					215					220				
Asn	Thr	Tyr	Gln	Thr	Ala	Phe	Arg	Gly	Leu	Asn	Thr	Arg	Leu	His	Asp
225					230				235					240	
Met	Leu	Glu	Phe	Arg	Thr	Tyr	Met	Phe	Leu	Asn	Val	Phe	Glu	Tyr	Val
			245					250						255	
Ser	Ile	Trp	Ser	Leu	Phe	Lys	Tyr	Gln	Ser	Leu	Met	Val	Ser	Ser	Gly
			260					265					270		
Ala	Asn	Leu	Tyr	Ala	Ser	Gly	Ser	Gly	Pro	Gln	Gln	Thr	Gln	Ser	Phe
		275					280					285			
Thr	Ala	Gln	Asn	Trp	Pro	Phe	Leu	Tyr	Ser	Leu	Phe	Gln	Val	Asn	Ser
	290					295					300				
Asn	Tyr	Ile	Leu	Ser	Gly	Ile	Ser	Gly	Thr	Arg	Leu	Ser	Ile	Thr	Phe
305					310					315				320	
Pro	Asn	Ile	Gly	Gly	Leu	Pro	Gly	Ser	Thr	Thr	Thr	His	Ser	Leu	Asn
			325					330						335	
Ser	Ala	Arg	Val	Asn	Tyr	Ser	Gly	Gly	Val	Ser	Ser	Gly	Leu	Ile	Gly
			340					345					350		
Ala	Thr	Asn	Leu	Asn	His	Asn	Phe	Asn	Cys	Ser	Thr	Val	Leu	Pro	Pro
		355					360					365			
Leu	Ser	Thr	Pro	Phe	Val	Arg	Ser	Trp	Leu	Asp	Ser	Gly	Thr	Asp	Arg
	370					375					380				
Glu	Gly	Val	Ala	Thr	Ser	Thr	Asn	Trp	Gln	Thr	Glu	Ser	Phe	Gln	Thr
385					390				395					400	
Thr	Leu	Ser	Leu	Arg	Cys	Gly	Ala	Phe	Ser	Ala	Arg	Gly	Asn	Ser	Asn
				405					410					415	
Tyr	Phe	Pro	Asp	Tyr	Phe	Ile	Arg	Asn	Ile	Ser	Gly	Val	Pro	Leu	Val
			420					425					430		
Ile	Arg	Asn	Glu	Asp	Leu	Thr	Arg	Pro	Leu	His	Tyr	Asn	Gln	Ile	Arg
		435					440					445			
Asn	Ile	Glu	Ser	Pro	Ser	Gly	Thr	Pro	Gly	Gly	Ala	Arg	Ala	Tyr	Leu
	450					455					460				
Val	Ser	Val	His	Asn	Arg	Lys	Asn	Asn	Ile	Tyr	Ala	Ala	Asn	Glu	Asn
465					470				475					480	
Gly	Thr	Met	Ile	His	Leu	Ala	Pro	Glu	Asp	Tyr	Thr	Gly	Phe	Thr	Ile
				485					490					495	
Ser	Pro	Ile	His	Ala	Thr	Gln	Val	Asn	Asn	Gln	Thr	Arg	Thr	Phe	Ile
			500					505					510		
Ser	Glu	Lys	Phe	Gly	Asn	Gln	Gly	Asp	Ser	Leu	Arg	Phe	Glu	Gln	Ser
		515					520					525			
Asn	Thr	Thr	Ala	Arg	Tyr	Thr	Leu	Arg	Gly	Asn	Gly	Asn	Ser	Tyr	Asn
	530					535					540				
Leu	Tyr	Leu	Arg	Val	Ser	Ser	Ile	Gly	Asn	Ser	Thr	Ile	Arg	Val	Thr
545					550				555					560	
Ile	Asn	Gly	Arg	Val	Tyr	Thr	Val	Ser	Asn	Val	Asn	Thr	Thr	Thr	Asn
				565					570					575	
Asn	Asp	Gly	Val	Asn	Asp	Asn	Gly	Ala	Arg	Phe	Ser	Asp	Ile	Asn	Ile
			580					585					590		
Gly	Asn	Ile	Val	Ala	Ser	Asp	Asn	Thr	Asn	Val	Thr	Leu	Asp	Ile	Asn

	595					600						605					
Val	Thr	Leu	Asn	Ser	Gly	Thr	Pro	Phe	Asp	Leu	Met	Asn	Ile	Met	Phe		
	610					615					620						
Val	Pro	Thr	Asn	Leu	Pro	Pro	Leu	Tyr									
625					630												

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 <212> PRT
 <213> *Bacillus thuringiensis*

<400> 12

Met	Ile	Arg	Lys	Gly	Gly	Arg	Lys	Met	Asn	Pro	Asn	Asn	Arg	Ser	Glu		
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His	Asp	Thr	Ile	Lys	Thr	Thr	Glu	Asn	Asn	Glu	Val	Pro	Thr	Asn	His		
			20				25						30				
Val	Gln	Tyr	Pro	Leu	Ala	Glu	Thr	Pro	Asn	Pro	Thr	Leu	Glu	Asp	Leu		
	35					40						45					
Asn	Tyr	Lys	Glu	Phe	Leu	Arg	Met	Thr	Ala	Asp	Asn	Asn	Thr	Glu	Ala		
	50					55					60						
Leu	Asp	Ser	Ser	Thr	Thr	Lys	Asp	Val	Ile	Gln	Lys	Gly	Ile	Ser	Val		
65				70						75					80		
Val	Gly	Asp	Leu	Leu	Gly	Val	Val	Gly	Phe	Pro	Phe	Gly	Gly	Ala	Leu		
			85					90						95			
Val	Ser	Phe	Tyr	Thr	Asn	Phe	Leu	Asn	Thr	Ile	Trp	Pro	Ser	Glu	Asp		
			100					105					110				
Pro	Trp	Lys	Ala	Phe	Met	Glu	Gln	Val	Glu	Ala	Leu	Met	Asp	Gln	Lys		
	115					120						125					
Ile	Ala	Asp	Tyr	Ala	Lys	Asn	Lys	Ala	Leu	Ala	Glu	Leu	Gln	Gly	Leu		
	130					135					140						
Gln	Asn	Asn	Val	Glu	Asp	Tyr	Val	Ser	Ala	Leu	Ser	Ser	Trp	Gln	Lys		
145				150						155					160		
Asn	Pro	Val	Ser	Ser	Arg	Asn	Pro	His	Ser	Gln	Gly	Arg	Ile	Arg	Glu		
			165					170						175			
Leu	Phe	Ser	Gln	Ala	Glu	Ser	His	Phe	Arg	Asn	Ser	Met	Pro	Ser	Phe		
			180					185					190				
Ala	Ile	Ser	Gly	Tyr	Glu	Val	Leu	Phe	Leu	Thr	Thr	Tyr	Ala	Gln	Ala		
	195						200					205					
Ala	Asn	Thr	His	Leu	Phe	Leu	Leu	Lys	Asp	Ala	Gln	Ile	Tyr	Gly	Glu		
	210					215					220						
Glu	Trp	Gly	Tyr	Glu	Lys	Glu	Asp	Ile	Ala	Glu	Phe	Tyr	Lys	Arg	Gln		
225				230						235					240		
Leu	Lys	Leu	Thr	Gln	Glu	Tyr	Thr	Asp	His	Cys	Val	Lys	Trp	Tyr	Asn		
			245					250						255			
Val	Gly	Leu	Asp	Lys	Leu	Arg	Gly	Ser	Ser	Tyr	Glu	Ser	Trp	Val	Asn		
			260					265					270				
Phe	Asn	Arg	Tyr	Arg	Arg	Glu	Met	Thr	Leu	Thr	Val	Leu	Asp	Leu	Ile		
	275					280						285					
Ala	Leu	Phe	Pro	Leu	Tyr	Asp	Val	Arg	Leu	Tyr	Pro	Lys	Glu	Val	Lys		
	290					295					300						
Thr	Glu	Leu	Thr	Arg	Asp	Val	Leu	Thr	Asp	Pro	Ile	Val	Gly	Val	Asn		
305				310						315					320		
Asn	Leu	Arg	Gly	Tyr	Gly	Thr	Thr	Phe	Ser	Asn	Ile	Glu	Asn	Tyr	Ile		
			325					330						335			
Arg	Lys	Pro	His	Leu	Phe	Asp	Tyr	Leu	His	Arg	Ile	Gln	Phe	His	Thr		
			340					345					350				

Arg	Phe	Gln	Pro	Gly	Tyr	Tyr	Gly	Asn	Asp	Ser	Phe	Asn	Tyr	Trp	Ser
		355					360					365			
Gly	Asn	Tyr	Val	Ser	Thr	Arg	Pro	Ser	Ile	Gly	Ser	Asn	Asp	Ile	Ile
		370					375				380				
Thr	Ser	Pro	Phe	Tyr	Gly	Asn	Lys	Ser	Ser	Glu	Pro	Val	Gln	Asn	Leu
385					390					395					400
Glu	Phe	Asn	Gly	Glu	Lys	Val	Tyr	Arg	Ala	Val	Ala	Asn	Thr	Asn	Leu
			405						410					415	
Ala	Val	Trp	Pro	Ser	Ala	Val	Tyr	Ser	Gly	Val	Thr	Lys	Val	Glu	Phe
			420					425					430		
Ser	Gln	Tyr	Asn	Asp	Gln	Thr	Asp	Glu	Ala	Ser	Thr	Gln	Thr	Tyr	Asp
		435					440					445			
Ser	Lys	Arg	Asn	Val	Gly	Ala	Val	Ser	Trp	Asp	Ser	Ile	Asp	Gln	Leu
		450				455					460				
Pro	Pro	Glu	Thr	Thr	Asp	Glu	Pro	Leu	Glu	Lys	Gly	Tyr	Ser	His	Gln
465					470					475					480
Leu	Asn	Tyr	Val	Met	Cys	Phe	Leu	Met	Gln	Gly	Ser	Arg	Gly	Thr	Ile
				485					490					495	
Pro	Val	Leu	Thr	Trp	Thr	His	Lys	Ser	Val	Asp	Phe	Phe	Asn	Met	Ile
			500					505					510		
Asp	Ser	Lys	Lys	Ile	Thr	Gln	Leu	Pro	Leu	Val	Lys	Ala	Tyr	Lys	Leu
		515					520					525			
Gln	Ser	Gly	Ala	Ser	Val	Val	Ala	Gly	Pro	Arg	Phe	Thr	Gly	Gly	Asp
		530				535					540				
Ile	Ile	Gln	Cys	Thr	Glu	Asn	Gly	Ser	Ala	Ala	Thr	Ile	Tyr	Val	Thr
545					550					555					560
Pro	Asp	Val	Ser	Tyr	Ser	Gln	Lys	Tyr	Arg	Ala	Arg	Ile	His	Tyr	Ala
				565					570					575	
Ser	Thr	Ser	Gln	Ile	Thr	Phe	Thr	Leu	Ser	Leu	Asp	Gly	Ala	Pro	Phe
			580					585					590		
Asn	Gln	Tyr	Tyr	Phe	Asp	Lys	Thr	Ile	Asn	Lys	Gly	Asp	Thr	Leu	Thr
		595					600					605			
Tyr	Asn	Ser	Phe	Asn	Leu	Ala	Ser	Phe	Ser	Thr	Pro	Phe	Glu	Leu	Ser
		610				615					620				
Gly	Asn	Asn	Leu	Gln	Ile	Gly	Val	Thr	Gly	Leu	Ser	Ala	Gly	Asp	Lys
625					630					635					640
Val	Tyr	Ile	Asp	Lys	Ile	Glu	Phe	Ile	Pro	Val	Asn				
				645					650						

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 <213> Bacillus thuringiensis

<400> 13

Met	Asn	Pro	Asn	Asn	Arg	Ser	Glu	His	Asp	Thr	Ile	Lys	Val	Thr	Pro
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			20					25					30		
Pro	Asn	Ser	Thr	Leu	Glu	Glu	Leu	Asn	Tyr	Lys	Glu	Phe	Leu	Arg	Met
		35					40					45			
Thr	Glu	Asp	Ser	Ser	Thr	Glu	Val	Leu	Asp	Asn	Ser	Thr	Val	Lys	Asp
		50				55					60				
Ala	Val	Gly	Thr	Gly	Ile	Ser	Val	Val	Gly	Gln	Ile	Leu	Gly	Val	Val
65					70					75				80	
Gly	Val	Pro	Phe	Ala	Gly	Ala	Leu	Thr	Ser	Phe	Tyr	Gln	Ser	Phe	Leu

				85				90				95			
Asn	Thr	Ile	Trp	Pro	Ser	Asp	Ala	Asp	Pro	Trp	Lys	Ala	Phe	Met	Ala
100				105				110							
Gln	Val	Glu	Val	Leu	Ile	Asp	Lys	Lys	Ile	Glu	Glu	Tyr	Ala	Lys	Ser
115				120				125							
Lys	Ala	Leu	Ala	Glu	Leu	Gln	Gly	Leu	Gln	Asn	Asn	Phe	Glu	Asp	Tyr
130				135				140							
Val	Asn	Ala	Leu	Asn	Ser	Trp	Lys	Lys	Thr	Pro	Leu	Ser	Leu	Arg	Ser
145	150				155				160						
Lys	Arg	Ser	Gln	Asp	Arg	Ile	Arg	Glu	Leu	Phe	Ser	Gln	Ala	Glu	Ser
165				170				175							
His	Phe	Arg	Asn	Ser	Met	Pro	Ser	Phe	Ala	Val	Ser	Lys	Phe	Glu	Val
180				185				190							
Leu	Phe	Leu	Pro	Thr	Tyr	Ala	Gln	Ala	Ala	Asn	Thr	His	Leu	Leu	Leu
195				200				205							
Leu	Lys	Asp	Ala	Gln	Val	Phe	Gly	Glu	Glu	Trp	Gly	Tyr	Ser	Ser	Glu
210				215				220							
Asp	Val	Ala	Glu	Phe	Tyr	His	Arg	Gln	Leu	Lys	Leu	Thr	Gln	Gln	Tyr
225	230				235				240						
Thr	Asp	His	Cys	Val	Asn	Trp	Tyr	Asn	Val	Gly	Leu	Asn	Gly	Leu	Arg
245				250				255							
Gly	Ser	Thr	Tyr	Asp	Ala	Trp	Val	Lys	Phe	Asn	Arg	Phe	Arg	Arg	Glu
260				265				270							
Met	Thr	Leu	Thr	Val	Leu	Asp	Leu	Ile	Val	Leu	Phe	Pro	Phe	Tyr	Asp
275				280				285							
Ile	Arg	Leu	Tyr	Ser	Lys	Gly	Val	Lys	Thr	Glu	Leu	Thr	Arg	Asp	Ile
290				295				300							
Phe	Thr	Asp	Pro	Ile	Phe	Ser	Leu	Asn	Thr	Leu	Gln	Glu	Tyr	Gly	Pro
305	310				315				320						
Thr	Phe	Leu	Ser	Ile	Glu	Asn	Ser	Ile	Arg	Lys	Pro	His	Leu	Phe	Asp
325				330				335							
Tyr	Leu	Gln	Gly	Ile	Glu	Phe	His	Thr	Arg	Leu	Gln	Pro	Gly	Tyr	Phe
340				345				350							
Gly	Lys	Asp	Ser	Phe	Asn	Tyr	Trp	Ser	Gly	Asn	Tyr	Val	Glu	Thr	Arg
355				360				365							
Pro	Ser	Ile	Gly	Ser	Ser	Lys	Thr	Ile	Thr	Ser	Pro	Phe	Tyr	Gly	Asp
370				375				380							
Lys	Ser	Thr	Glu	Pro	Val	Gln	Lys	Leu	Ser	Phe	Asp	Gly	Gln	Lys	Val
385	390				395				400						
Tyr	Arg	Thr	Ile	Ala	Asn	Thr	Asp	Val	Ala	Ala	Trp	Pro	Asn	Gly	Lys
405				410				415							
Val	Tyr	Leu	Gly	Val	Thr	Lys	Val	Asp	Phe	Ser	Gln	Tyr	Asp	Asp	Gln
420				425				430							
Lys	Asn	Glu	Thr	Ser	Thr	Gln	Thr	Tyr	Asp	Ser	Lys	Arg	Asn	Asn	Gly
435				440				445							
His	Val	Ser	Ala	Gln	Asp	Ser	Ile	Asp	Gln	Leu	Pro	Pro	Glu	Thr	Thr
450				455				460							
Asp	Glu	Pro	Leu	Glu	Lys	Ala	Tyr	Ser	His	Gln	Leu	Asn	Tyr	Ala	Glu
465	470				475				480						
Cys	Phe	Leu	Met	Gln	Asp	Arg	Arg	Gly	Thr	Ile	Pro	Phe	Phe	Thr	Trp
485				490				495							
Thr	His	Arg	Ser	Val	Asp	Phe	Phe	Asn	Thr	Ile	Asp	Ala	Glu	Lys	Ile
500				505				510							
Thr	Gln	Leu	Pro	Val	Val	Lys	Ala	Tyr	Ala	Leu	Ser	Ser	Gly	Ala	Ser
515				520				525							
Ile	Ile	Glu	Gly	Pro	Gly	Phe	Thr	Gly	Gly	Asn	Leu	Phe	Leu	Lys	
530				535				540							

Glu	Ser	Ser	Asn	Ser	Ile	Ala	Lys	Phe	Lys	Val	Thr	Leu	Asn	Ser	Ala
545					550					555					560
Ala	Leu	Leu	Gln	Arg	Tyr	Arg	Val	Arg	Ile	Arg	Tyr	Ala	Ser	Thr	Thr
				565					570						575
Asn	Leu	Arg	Leu	Phe	Val	Gln	Asn	Ser	Asn	Asn	Asp	Phe	Leu	Val	Ile
			580					585					590		
Tyr	Ile	Asn	Lys	Thr	Met	Asn	Lys	Asp	Asp	Asp	Leu	Thr	Tyr	Gln	Thr
		595					600					605			
Phe	Asp	Leu	Ala	Thr	Thr	Asn	Ser	Asn	Met	Gly	Phe	Ser	Gly	Asp	Lys
	610					615					620				
Asn	Glu	Leu	Ile	Ile	Gly	Ala	Glu	Ser	Phe	Val	Ser	Asn	Glu	Lys	Ile
625					630					635					640
Tyr	Ile	Asp	Lys	Ile	Glu	Phe	Ile	Pro	Val	Gln	Leu				
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<210> 14

<211> 1180

<212> PRT

<213> *Bacillus thuringiensis*

<400> 14

Met	Asn	Pro	Tyr	Gln	Asn	Lys	Asn	Glu	Tyr	Glu	Thr	Leu	Asn	Ala	Ser
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Gln	Lys	Lys	Leu	Asn	Ile	Ser	Asn	Asn	Tyr	Thr	Arg	Tyr	Pro	Ile	Glu
			20					25					30		
Asn	Ser	Pro	Lys	Gln	Leu	Leu	Gln	Ser	Thr	Asn	Tyr	Lys	Asp	Trp	Leu
		35					40					45			
Asn	Met	Cys	Gln	Gln	Asn	Gln	Gln	Tyr	Gly	Gly	Asp	Phe	Glu	Thr	Phe
	50					55					60				
Ile	Asp	Ser	Gly	Glu	Leu	Ser	Ala	Tyr	Thr	Ile	Val	Val	Gly	Thr	Val
65					70					75					80
Leu	Thr	Gly	Phe	Gly	Phe	Thr	Thr	Pro	Leu	Gly	Leu	Ala	Leu	Ile	Gly
				85				90						95	
Phe	Gly	Thr	Leu	Ile	Pro	Val	Leu	Phe	Pro	Ala	Gln	Asp	Gln	Ser	Asn
			100					105					110		
Thr	Trp	Ser	Asp	Phe	Ile	Thr	Gln	Thr	Lys	Asn	Ile	Ile	Lys	Lys	Glu
		115					120					125			
Ile	Ala	Ser	Thr	Tyr	Ile	Ser	Asn	Ala	Asn	Lys	Ile	Leu	Asn	Arg	Ser
	130					135					140				
Phe	Asn	Val	Ile	Ser	Thr	Tyr	His	Asn	His	Leu	Lys	Thr	Trp	Glu	Asn
145					150					155					160
Asn	Pro	Asn	Pro	Gln	Asn	Thr	Gln	Asp	Val	Arg	Thr	Gln	Ile	Gln	Leu
				165					170					175	
Val	His	Tyr	His	Phe	Gln	Asn	Val	Ile	Pro	Glu	Leu	Val	Asn	Ser	Cys
			180					185					190		
Pro	Pro	Asn	Pro	Ser	Asp	Cys	Asp	Tyr	Tyr	Asn	Ile	Leu	Val	Leu	Ser
		195					200					205			
Ser	Tyr	Ala	Gln	Ala	Ala	Asn	Leu	His	Leu	Thr	Val	Leu	Asn	Gln	Ala
	210					215					220				
Val	Lys	Phe	Glu	Ala	Tyr	Leu	Lys	Asn	Asn	Arg	Gln	Phe	Asp	Tyr	Leu
225					230					235					240
Glu	Pro	Leu	Pro	Thr	Ala	Ile	Asp	Tyr	Tyr	Pro	Val	Leu	Thr	Lys	Ala
				245					250					255	
Ile	Glu	Asp	Tyr	Thr	Asn	Tyr	Cys	Val	Thr	Thr	Tyr	Lys	Lys	Gly	Leu
		260						265					270		
Asn	Leu	Ile	Lys	Thr	Thr	Pro	Asp	Ser	Asn	Leu	Asp	Gly	Asn	Ile	Asn

		275					280					285					
Trp	Asn	Thr	Tyr	Asn	Thr	Tyr	Arg	Thr	Lys	Met	Thr	Thr	Ala	Val	Leu		
	290						295					300					
Asp	Leu	Val	Ala	Leu	Phe	Pro	Asn	Tyr	Asp	Val	Gly	Lys	Tyr	Pro	Ile		
305					310					315					320		
Gly	Val	Gln	Ser	Glu	Leu	Thr	Arg	Glu	Ile	Tyr	Gln	Val	Leu	Asn	Phe		
				325					330					335			
Glu	Glu	Ser	Pro	Tyr	Lys	Tyr	Tyr	Asp	Phe	Gln	Tyr	Gln	Glu	Asp	Ser		
			340					345					350				
Leu	Thr	Arg	Arg	Pro	His	Leu	Phe	Thr	Trp	Leu	Asp	Ser	Leu	Asn	Phe		
		355					360					365					
Tyr	Glu	Lys	Ala	Gln	Thr	Thr	Pro	Asn	Asn	Phe	Phe	Thr	Ser	His	Tyr		
	370					375					380						
Asn	Met	Phe	His	Tyr	Thr	Leu	Asp	Asn	Ile	Ser	Gln	Lys	Ser	Ser	Val		
385					390					395					400		
Phe	Gly	Asn	His	Asn	Val	Thr	Asp	Lys	Leu	Lys	Ser	Leu	Gly	Leu	Ala		
				405					410					415			
Thr	Asn	Ile	Tyr	Ile	Phe	Leu	Leu	Asn	Val	Ile	Ser	Leu	Asp	Asn	Lys		
		420						425					430				
Tyr	Leu	Asn	Asp	Tyr	Asn	Asn	Ile	Ser	Lys	Met	Asp	Phe	Phe	Ile	Thr		
	435					440					445						
Asn	Gly	Thr	Arg	Leu	Leu	Glu	Lys	Glu	Leu	Thr	Ala	Gly	Ser	Gly	Gln		
	450					455					460						
Ile	Thr	Tyr	Asp	Val	Asn	Lys	Asn	Ile	Phe	Gly	Leu	Pro	Ile	Leu	Lys		
465					470					475					480		
Arg	Arg	Glu	Asn	Gln	Gly	Asn	Pro	Thr	Leu	Phe	Pro	Thr	Tyr	Asp	Asn		
			485					490					495				
Tyr	Ser	His	Ile	Leu	Ser	Phe	Ile	Lys	Ser	Leu	Ser	Ile	Pro	Ala	Thr		
		500						505					510				
Tyr	Lys	Thr	Gln	Val	Tyr	Thr	Phe	Ala	Trp	Thr	His	Ser	Ser	Val	Asp		
	515						520					525					
Pro	Lys	Asn	Thr	Ile	Tyr	Thr	His	Leu	Thr	Thr	Gln	Ile	Pro	Ala	Val		
	530					535					540						
Lys	Ala	Asn	Ser	Leu	Gly	Thr	Ala	Ser	Lys	Val	Val	Gln	Gly	Pro	Gly		
545					550					555					560		
His	Thr	Gly	Gly	Asp	Leu	Ile	Asp	Phe	Lys	Asp	His	Phe	Lys	Ile	Thr		
				565					570					575			
Cys	Gln	His	Ser	Asn	Phe	Gln	Gln	Ser	Tyr	Phe	Ile	Arg	Ile	Arg	Tyr		
			580					585				590					
Ala	Ser	Asn	Gly	Ser	Ala	Asn	Thr	Arg	Ala	Val	Ile	Asn	Leu	Ser	Ile		
	595						600				605						
Pro	Gly	Val	Ala	Glu	Leu	Gly	Met	Ala	Leu	Asn	Pro	Thr	Phe	Ser	Gly		
	610					615					620						
Thr	Asp	Tyr	Thr	Asn	Leu	Lys	Tyr	Lys	Asp	Phe	Gln	Tyr	Leu	Glu	Phe		
625					630					635					640		
Ser	Asn	Glu	Val	Lys	Phe	Ala	Pro	Asn	Gln	Asn	Ile	Ser	Leu	Val	Phe		
				645					650					655			
Asn	Arg	Ser	Asp	Val	Tyr	Thr	Asn	Thr	Thr	Val	Leu	Ile	Asp	Lys	Ile		
		660						665					670				
Glu	Phe	Leu	Pro	Ile	Thr	Arg	Ser	Ile	Arg	Glu	Asp	Arg	Glu	Lys	Gln		
	675						680				685						
Lys	Leu	Glu	Thr	Val	Gln	Gln	Ile	Ile	Asn	Thr	Phe	Tyr	Ala	Asn	Pro		
	690					695					700						
Ile	Lys	Asn	Thr	Leu	Gln	Ser	Glu	Leu	Thr	Asp	Tyr	Asp	Ile	Asp	Gln		
705					710					715					720		
Ala	Ala	Asn	Leu	Val	Glu	Cys	Ile	Ser	Glu	Glu	Leu	Tyr	Pro	Lys	Glu		
				725					730					735			

Lys	Met	Leu	Leu	Leu	Asp	Glu	Val	Lys	Asn	Ala	Lys	Gln	Leu	Ser	Gln	
		740						745				750				
Ser	Arg	Asn	Val	Leu	Gln	Asn	Gly	Asp	Phe	Glu	Ser	Ala	Thr	Leu	Gly	
		755					760					765				
Trp	Thr	Thr	Ser	Asp	Asn	Ile	Thr	Ile	Gln	Glu	Asp	Asp	Pro	Ile	Phe	
	770					775					780					
Lys	Gly	His	Tyr	Leu	His	Met	Ser	Gly	Ala	Arg	Asp	Ile	Asp	Gly	Thr	
785					790					795					800	
Ile	Phe	Pro	Thr	Tyr	Ile	Phe	Gln	Lys	Ile	Asp	Glu	Ser	Lys	Leu	Lys	
				805					810					815		
Pro	Tyr	Thr	Arg	Tyr	Leu	Val	Arg	Gly	Phe	Val	Gly	Ser	Ser	Lys	Asp	
			820					825					830			
Val	Glu	Leu	Val	Val	Ser	Arg	Tyr	Gly	Glu	Glu	Ile	Asp	Ala	Ile	Met	
		835					840					845				
Asn	Val	Pro	Ala	Asp	Leu	Asn	Tyr	Leu	Tyr	Pro	Ser	Thr	Phe	Asp	Cys	
	850					855					860					
Glu	Gly	Ser	Asn	Arg	Cys	Glu	Thr	Ser	Ala	Val	Pro	Ala	Asn	Ile	Gly	
865				870						875					880	
Asn	Thr	Ser	Asp	Met	Leu	Tyr	Ser	Cys	Gln	Tyr	Asp	Thr	Gly	Lys	Lys	
				885					890					895		
His	Val	Val	Cys	Gln	Asp	Ser	His	Gln	Phe	Ser	Phe	Thr	Ile	Asp	Thr	
			900					905					910			
Gly	Ala	Leu	Asp	Thr	Asn	Glu	Asn	Ile	Gly	Val	Trp	Val	Met	Phe	Lys	
		915				920						925				
Ile	Ser	Ser	Pro	Asp	Gly	Tyr	Ala	Ser	Leu	Asp	Asn	Leu	Glu	Val	Ile	
	930					935					940					
Glu	Glu	Gly	Pro	Ile	Asp	Gly	Glu	Ala	Leu	Ser	Arg	Val	Lys	His	Met	
945					950					955					960	
Glu	Lys	Lys	Trp	Asn	Asp	Gln	Met	Glu	Ala	Lys	Arg	Ser	Glu	Thr	Gln	
				965					970					975		
Gln	Ala	Tyr	Asp	Val	Ala	Lys	Gln	Ala	Ile	Asp	Ala	Leu	Phe	Thr	Asn	
			980					985					990			
Val	Gln	Asp	Glu	Ala	Leu	Gln	Phe	Asp	Thr	Thr	Leu	Ala	Gln	Ile	Gln	
		995					1000						1005			
Tyr	Ala	Glu	Tyr	Leu	Val	Gln	Ser	Ile	Pro	Tyr	Val	Tyr	Asn	Asp	Trp	
	1010					1015					1020					
Leu	Ser	Asp	Val	Pro	Gly	Met	Asn	Tyr	Asp	Ile	Tyr	Val	Glu	Leu	Asp	
1025					1030					1035					1040	
Ala	Arg	Val	Ala	Gln	Ala	Arg	Tyr	Leu	Tyr	Asp	Thr	Arg	Asn	Ile	Ile	
				1045					1050					1055		
Lys	Asn	Gly	Asp	Phe	Thr	Gln	Gly	Val	Met	Gly	Trp	His	Val	Thr	Gly	
			1060					1065					1070			
Asn	Ala	Asp	Val	Gln	Gln	Ile	Asp	Gly	Val	Ser	Val	Leu	Val	Leu	Ser	
		1075					1080					1085				
Asn	Trp	Ser	Ala	Gly	Val	Ser	Gln	Asn	Val	His	Leu	Gln	His	Asn	His	
	1090					1095						1100				
Gly	Tyr	Val	Leu	Arg	Val	Ile	Ala	Lys	Lys	Glu	Gly	Pro	Gly	Asn	Gly	
1105					1110					1115					1120	
Tyr	Val	Thr	Leu	Met	Asp	Cys	Glu	Glu	Asn	Gln	Glu	Lys	Leu	Thr	Phe	
				1125					1130					1135		
Thr	Ser	Cys	Glu	Gly	Tyr	Ile	Thr	Lys	Thr	Val	Asp	Val	Phe	Pro		
			1140				1145					1150				
Asp	Thr	Asp	Arg	Val	Arg	Ile	Glu	Ile	Gly	Glu	Thr	Glu	Gly	Ser	Phe	
		1155					1160					1165				
Tyr	Ile	Glu	Ser	Ile	Glu	Leu	Ile	Cys	Met	Asn	Glu					
	1170					1175					1180					

<210> 15
 <211> 1136
 <212> PRT
 <213> *Bacillus thuringiensis*

<400> 15
 Met Asn Ser Gly Tyr Pro Leu Ala Asn Asp Leu Gln Gly Ser Met Lys
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 Asn Thr Asn Tyr Lys Asp Trp Leu Ala Met Cys Glu Asn Asn Gln Gln
 20 25 30
 Tyr Gly Val Asn Pro Ala Ala Ile Asn Ser Ser Ser Val Ser Thr Ala
 35 40 45
 Leu Lys Val Ala Gly Ala Ile Leu Lys Phe Val Asn Pro Pro Ala Gly
 50 55 60
 Thr Val Leu Thr Val Leu Ser Ala Val Leu Pro Ile Leu Trp Pro Thr
 65 70 75 80
 Asn Thr Pro Thr Pro Glu Arg Val Trp Asn Asp Phe Met Thr Asn Thr
 85 90 95
 Gly Asn Leu Ile Asp Gln Thr Val Thr Ala Tyr Val Arg Thr Asp Ala
 100 105 110
 Asn Ala Lys Met Thr Val Val Lys Asp Tyr Leu Asp Gln Tyr Thr Thr
 115 120 125
 Lys Phe Asn Thr Trp Lys Arg Glu Pro Asn Asn Gln Ser Tyr Arg Thr
 130 135 140
 Ala Val Ile Thr Gln Phe Asn Leu Thr Ser Ala Lys Leu Arg Glu Thr
 145 150 155 160
 Ala Val Tyr Phe Ser Asn Leu Val Gly Tyr Glu Leu Leu Leu Leu Pro
 165 170 175
 Ile Tyr Ala Gln Val Ala Asn Phe Asn Leu Leu Leu Ile Arg Asp Gly
 180 185 190
 Leu Ile Asn Ala Gln Glu Trp Ser Leu Ala Arg Ser Ala Gly Asp Gln
 195 200 205
 Leu Tyr Asn Thr Met Val Gln Tyr Thr Lys Glu Tyr Ile Ala His Ser
 210 215 220
 Ile Thr Trp Tyr Asn Lys Gly Leu Asp Val Leu Arg Asn Lys Ser Asn
 225 230 235 240
 Gly Gln Trp Ile Thr Phe Asn Asp Tyr Lys Arg Glu Met Thr Ile Gln
 245 250 255
 Val Leu Asp Ile Leu Ala Leu Phe Ala Ser Tyr Asp Pro Arg Arg Tyr
 260 265 270
 Pro Ala Asp Lys Ile Asp Asn Thr Lys Leu Ser Lys Thr Glu Phe Thr
 275 280 285
 Arg Glu Ile Tyr Thr Ala Leu Val Glu Ser Pro Ser Ser Lys Ser Ile
 290 295 300
 Ala Ala Leu Glu Ala Ala Leu Thr Arg Asp Val His Leu Phe Thr Trp
 305 310 315 320
 Leu Lys Arg Val Asp Phe Trp Thr Asn Thr Ile Tyr Gln Asp Leu Arg
 325 330 335
 Phe Leu Ser Ala Asn Lys Ile Gly Phe Ser Tyr Thr Asn Ser Ser Ala
 340 345 350
 Met Gln Glu Ser Gly Ile Tyr Gly Ser Ser Gly Phe Gly Ser Asn Leu
 355 360 365
 Thr His Gln Ile Gln Leu Asn Ser Asn Val Tyr Lys Thr Ser Ile Thr
 370 375 380
 Asp Thr Ser Ser Pro Ser Asn Arg Val Thr Lys Met Asp Phe Tyr Lys
 385 390 395 400

850		855		860	
Thr Ile Asp Thr Gly Ala Leu Asp Thr Asn Glu Asn Ile Gly Val Trp					
865		870		875	880
Val Met Phe Lys Ile Ser Ser Pro Asp Gly Tyr Ala Ser Leu Asp Asn					
	885		890		895
Leu Glu Val Ile Glu Glu Gly Pro Ile Asp Gly Glu Ala Leu Ser Arg					
	900		905		910
Val Lys His Met Glu Lys Lys Trp Asn Asp Gln Met Glu Ala Lys Arg					
	915		920		925
Ser Glu Thr Gln Gln Ala Tyr Asp Val Ala Lys Gln Ala Ile Asp Ala					
	930		935		940
Leu Phe Thr Asn Val Gln Asp Glu Ala Leu Gln Phe Asp Thr Thr Leu					
945		950		955	960
Ala Gln Ile Gln Tyr Ala Glu Tyr Leu Val Gln Ser Ile Pro Tyr Val					
	965		970		975
Tyr Asn Asp Trp Leu Ser Asp Val Pro Gly Met Asn Tyr Asp Ile Tyr					
	980		985		990
Val Glu Leu Asp Ala Arg Val Ala Gln Ala Arg Tyr Leu Tyr Asp Thr					
	995		1000		1005
Arg Asn Ile Ile Lys Asn Gly Asp Phe Thr Gln Gly Val Met Gly Trp					
	1010		1015		1020
His Val Thr Gly Asn Ala Asp Val Gln Gln Ile Asp Gly Val Ser Val					
1025		1030		1035	1040
Leu Val Leu Ser Asn Trp Ser Ala Gly Val Ser Gln Asn Val His Leu					
	1045		1050		1055
Gln His Asn His Gly Tyr Val Leu Arg Val Ile Ala Lys Lys Glu Gly					
	1060		1065		1070
Pro Gly Asn Gly Tyr Val Thr Leu Met Asp Cys Glu Glu Asn Gln Glu					
	1075		1080		1085
Lys Leu Thr Phe Thr Ser Cys Glu Glu Gly Tyr Ile Thr Lys Thr Val					
	1090		1095		1100
Asp Val Phe Pro Asp Thr Asp Arg Val Arg Ile Glu Ile Gly Glu Thr					
1105		1110		1115	1120
Glu Gly Ser Phe Tyr Ile Glu Ser Ile Glu Leu Ile Cys Met Asn Glu					
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<210> 16

<211> 475

<212> PRT

<213> Bacillus thuringiensis

<400> 16

Met Ile Ile Asp Ser Lys Thr Thr Leu Pro Arg His Ser Leu Ile His					
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Thr Ile Lys Leu Asn Ser Asn Lys Lys Tyr Gly Pro Gly Asp Met Thr					
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Asn Gly Asn Gln Phe Ile Ile Ser Lys Gln Glu Trp Ala Thr Ile Gly					
	35		40		45
Ala Tyr Ile Gln Thr Gly Leu Gly Leu Pro Val Asn Glu Gln Gln Leu					
	50		55		60
Arg Thr His Val Asn Leu Ser Gln Asp Ile Ser Ile Pro Ser Asp Phe					
65		70		75	80
Ser Gln Leu Tyr Asp Val Tyr Cys Ser Asp Lys Thr Ser Ala Glu Trp					
	85		90		95
Trp Asn Lys Asn Leu Tyr Pro Leu Ile Ile Lys Ser Ala Asn Asp Ile					
	100		105		110

			20					25					30				
Lys	Asn	Met	Asn	Tyr	Gln	Asp	Phe	Leu	Ser	Ile	Thr	Glu	Arg	Glu	Gln		
		35					40					45					
Pro	Glu	Ala	Leu	Ala	Ser	Gly	Asn	Thr	Ala	Ile	Asn	Thr	Val	Val	Ser		
		50				55					60						
Val	Thr	Gly	Ala	Thr	Leu	Ser	Ala	Leu	Gly	Val	Pro	Gly	Ala	Ser	Phe		
65					70					75					80		
Ile	Thr	Asn	Phe	Tyr	Leu	Lys	Ile	Ala	Gly	Leu	Leu	Trp	Pro	Glu	Asn		
			85						90				95				
Gly	Lys	Ile	Trp	Asp	Glu	Phe	Met	Thr	Glu	Val	Glu	Ala	Leu	Ile	Asp		
			100					105					110				
Gln	Lys	Ile	Glu	Glu	Tyr	Val	Arg	Asn	Lys	Ala	Ile	Ala	Glu	Leu	Asp		
		115					120					125					
Gly	Leu	Gly	Ser	Ala	Leu	Asp	Lys	Tyr	Gln	Lys	Ala	Leu	Ala	Asp	Trp		
		130				135					140						
Leu	Gly	Lys	Gln	Asp	Asp	Pro	Glu	Ala	Ile	Leu	Ser	Val	Ala	Thr	Glu		
145				150						155					160		
Phe	Arg	Ile	Ile	Asp	Ser	Leu	Phe	Glu	Phe	Ser	Met	Pro	Ser	Phe	Lys		
				165					170					175			
Val	Thr	Gly	Tyr	Glu	Ile	Pro	Leu	Leu	Thr	Val	Tyr	Ala	Gln	Ala	Ala		
			180					185					190				
Asn	Leu	His	Leu	Ala	Leu	Leu	Arg	Asp	Ser	Thr	Leu	Tyr	Gly	Asp	Lys		
		195					200					205					
Trp	Gly	Phe	Thr	Gln	Asn	Asn	Ile	Glu	Glu	Asn	Tyr	Asn	Arg	Gln	Lys		
		210			215						220						
Lys	Arg	Ile	Ser	Glu	Tyr	Ser	Asp	His	Cys	Thr	Lys	Trp	Tyr	Asn	Ser		
225				230						235					240		
Gly	Leu	Ser	Arg	Leu	Asn	Gly	Ser	Thr	Tyr	Glu	Gln	Trp	Ile	Asn	Tyr		
			245						250					255			
Asn	Arg	Phe	Arg	Arg	Glu	Met	Ile	Leu	Met	Ala	Leu	Asp	Leu	Val	Ala		
		260						265					270				
Val	Phe	Pro	Phe	His	Asp	Pro	Arg	Arg	Tyr	Ser	Met	Glu	Thr	Ser	Thr		
		275					280					285					
Gln	Leu	Thr	Arg	Glu	Val	Tyr	Thr	Asp	Pro	Val	Ser	Leu	Ser	Ile	Ser		
		290				295					300						
Asn	Pro	Asp	Ile	Gly	Pro	Ser	Phe	Ser	Gln	Met	Glu	Asn	Thr	Ala	Ile		
305				310					315						320		
Arg	Thr	Pro	His	Leu	Val	Asp	Tyr	Leu	Asp	Glu	Leu	Tyr	Ile	Tyr	Thr		
			325					330					335				
Ser	Lys	Tyr	Lys	Ala	Phe	Ser	His	Glu	Ile	Gln	Pro	Asp	Leu	Phe	Tyr		
			340					345					350				
Trp	Ser	Ala	His	Lys	Val	Ser	Phe	Lys	Lys	Ser	Glu	Gln	Ser	Asn	Leu		
		355					360					365					
Tyr	Thr	Thr	Gly	Ile	Tyr	Gly	Lys	Thr	Ser	Gly	Tyr	Ile	Ser	Ser	Gly		
		370			375						380						
Ala	Tyr	Ser	Phe	His	Gly	Asn	Asp	Ile	Tyr	Arg	Thr	Leu	Ala	Ala	Pro		
385				390						395					400		
Ser	Val	Val	Val	Tyr	Pro	Tyr	Thr	Gln	Asn	Tyr	Gly	Val	Glu	Gln	Val		
			405						410					415			
Glu	Phe	Tyr	Gly	Val	Lys	Gly	His	Val	His	Tyr	Arg	Gly	Asp	Asn	Lys		
			420				425					430					
Tyr	Asp	Leu	Thr	Tyr	Asp	Ser	Ile	Asp	Gln	Leu	Pro	Pro	Asp	Gly	Glu		
		435				440						445					
Pro	Ile	His	Glu	Lys	Tyr	Thr	His	Arg	Leu	Cys	His	Ala	Thr	Ala	Ile		
		450				455					460						
Phe	Lys	Ser	Thr	Pro	Asp	Tyr	Asp	Asn	Ala	Thr	Ile	Pro	Ile	Phe	Ser		
465					470					475					480		

Trp	Thr	His	Arg	Ser	Ala	Glu	Tyr	Tyr	Asn	Arg	Ile	Tyr	Pro	Asn	Lys	
				485					490					495		
Ile	Thr	Lys	Ile	Pro	Ala	Val	Lys	Met	Tyr	Lys	Leu	Asp	Asp	Pro	Ser	
			500					505					510			
Thr	Val	Val	Lys	Gly	Pro	Gly	Phe	Thr	Gly	Gly	Asp	Leu	Val	Lys	Arg	
		515					520					525				
Gly	Ser	Thr	Gly	Tyr	Ile	Gly	Asp	Ile	Lys	Ala	Thr	Val	Asn	Ser	Pro	
	530					535					540					
Leu	Ser	Gln	Lys	Tyr	Arg	Val	Arg	Val	Arg	Tyr	Ala	Thr	Asn	Val	Ser	
545					550					555					560	
Gly	Gln	Phe	Asn	Val	Tyr	Ile	Asn	Asp	Lys	Ile	Thr	Leu	Gln	Thr	Lys	
			565						570						575	
Phe	Gln	Asn	Thr	Val	Glu	Thr	Ile	Gly	Glu	Gly	Lys	Asp	Leu	Thr	Tyr	
			580					585					590			
Gly	Ser	Phe	Gly	Tyr	Ile	Glu	Tyr	Ser	Thr	Thr	Ile	Gln	Phe	Pro	Asp	
		595					600					605				
Glu	His	Pro	Lys	Ile	Thr	Leu	His	Leu	Ser	Asp	Leu	Ser	Asn	Asn	Ser	
	610					615					620					
Ser	Phe	Tyr	Val	Asp	Ser	Ile	Glu	Phe	Ile	Pro	Val	Asp	Val	Asn	Tyr	
625					630					635					640	
Ala	Glu	Lys	Glu	Lys	Leu	Glu	Lys	Ala	Gln	Lys	Ala	Val	Asn	Thr	Leu	
				645					650					655		
Phe	Thr	Glu	Gly	Arg	Asn	Ala	Leu	Gln	Lys	Asp	Val	Thr	Asp	Tyr	Lys	
			660					665					670			
Val	Asp	Gln	Val	Ser	Ile	Leu	Val	Asp	Cys	Ile	Ser	Gly	Asp	Leu	Tyr	
		675					680					685				
Pro	Asn	Glu	Lys	Arg	Glu	Leu	Gln	Asn	Leu	Val	Lys	Tyr	Ala	Lys	Arg	
	690					695					700					
Leu	Ser	Tyr	Ser	Arg	Asn	Leu	Leu	Leu	Asp	Pro	Thr	Phe	Asp	Ser	Ile	
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Asn	Ser	Ser	Glu	Glu	Asn	Gly	Trp	Tyr	Gly	Ser	Asn	Gly	Ile	Val	Ile	
				725					730						735	
Gly	Asn	Gly	Asp	Phe	Val	Phe	Lys	Gly	Asn	Tyr	Leu	Ile	Phe	Ser	Gly	
			740					745					750			
Thr	Asn	Asp	Thr	Gln	Tyr	Pro	Thr	Tyr	Leu	Tyr	Gln	Lys	Ile	Asp	Glu	
		755						760					765			
Ser	Lys	Leu	Lys	Glu	Tyr	Thr	Arg	Tyr	Lys	Leu	Lys	Gly	Phe	Ile	Glu	
	770					775					780					
Ser	Ser	Gln	Asp	Leu	Glu	Ala	Tyr	Val	Ile	Arg	Tyr	Asp	Ala	Lys	His	
785					790					795					800	
Arg	Thr	Leu	Asp	Val	Ser	Asp	Asn	Leu	Leu	Pro	Asp	Ile	Leu	Pro	Glu	
				805					810					815		
Asn	Thr	Cys	Gly	Glu	Pro	Asn	Arg	Cys	Ala	Ala	Gln	Gln	Tyr	Leu	Asp	
			820					825					830			
Glu	Asn	Pro	Ser	Pro	Glu	Cys	Ser	Ser	Met	Gln	Asp	Gly	Ile	Leu	Ser	
		835					840					845				
Asp	Ser	His	Ser	Phe	Ser	Leu	Asn	Ile	Asp	Thr	Gly	Ser	Ile	Asn	His	
	850					855					860					
Asn	Glu	Asn	Leu	Gly	Ile	Trp	Val	Leu	Phe	Lys	Ile	Ser	Thr	Leu	Glu	
865					870					875					880	
Gly	Tyr	Ala	Lys	Phe	Gly	Asn	Leu	Glu	Val	Ile	Glu	Asp	Gly	Pro	Val	
				885					890					895		
Ile	Gly	Glu	Ala	Leu	Ala	Arg	Val	Lys	Arg	Gln	Glu	Thr	Lys	Trp	Arg	
			900					905					910			
Asn	Lys	Leu	Ala	Gln	Leu	Thr	Thr	Glu	Thr	Gln	Ala	Ile	Tyr	Thr	Arg	
		915						920				925				
Ala	Lys	Gln	Ala	Leu	Asp	Asn	Leu	Phe	Ala	Asn	Ala	Gln	Asp	Ser	His	

930	935	940
Leu Lys Arg Asp Val Thr Phe Ala Glu Ile Ala Ala Ala Arg Lys Ile		
945	950	955
Val Gln Ser Ile Arg Glu Ala Tyr Met Ser Trp Leu Ser Val Val Pro		960
	965	970
Gly Val Asn His Pro Ile Phe Thr Glu Leu Ser Gly Arg Val Gln Arg		975
	980	985
Ala Phe Gln Leu Tyr Asp Val Arg Asn Val Val Arg Asn Gly Arg Phe		990
	995	1000
Leu Asn Gly Leu Ser Asp Trp Ile Val Thr Ser Asp Val Lys Val Gln		1005
	1010	1015
Glu Glu Asn Gly Asn Asn Val Leu Val Leu Asn Asn Trp Asp Ala Gln		1020
1025	1030	1035
Val Leu Gln Asn Val Lys Leu Tyr Gln Asp Arg Gly Tyr Ile Leu His		1040
	1045	1050
Val Thr Ala Arg Lys Ile Gly Ile Gly Glu Gly Tyr Ile Thr Ile Thr		1055
	1060	1065
Asp Glu Glu Gly His Thr Asp Gln Leu Arg Phe Thr Ala Cys Glu Glu		1070
	1075	1080
Ile Asp Ala Ser Asn Ala Phe Ile Ser Gly Tyr Ile Thr Lys Glu Leu		1085
	1090	1095
Glu Phe Phe Pro Asp Thr Glu Lys Val His Ile Glu Ile Gly Glu Thr		1100
1105	1110	1115
Glu Gly Ile Phe Leu Val Glu Ser Ile Glu Leu Phe Leu Met Glu Glu		1120
	1125	1130
		1135
Leu Cys		

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35 40 45
Ser Gly Gly Glu Asn Pro Glu Leu Phe Gly Asn Pro Glu Thr Phe Ile
50 55 60
Ser Ser Ser Thr Ile Gln Thr Gly Ile Gly Ile Val Gly Arg Ile Leu
65 70 75 80
Gly Ala Leu Gly Val Pro Phe Ala Ser Gln Ile Ala Ser Phe Tyr Ser
85 90 95
Phe Ile Val Gly Gln Leu Trp Pro Ser Lys Ser Val Asp Ile Trp Gly
100 105 110
Glu Ile Met Glu Arg Val Glu Glu Val Asp Gln Lys Ile Glu Lys
115 120 125
Tyr Val Lys Asp Lys Ala Leu Ala Glu Leu Lys Gly Leu Gly Asn Ala
130 135 140
Leu Asp Val Tyr Gln Gln Ser Leu Glu Asp Trp Leu Glu Asn Arg Asn
145 150 155 160
Asp Ala Arg Thr Arg Ser Val Val Ser Asn Gln Phe Ile Ala Leu Asp
165 170 175

Leu	Asn	Phe	Val	Ser	Ser	Ile	Pro	Ser	Phe	Ala	Val	Ser	Gly	His	Glu
			180						185				190		
Val	Leu	Leu	Leu	Ala	Val	Tyr	Ala	Gln	Ala	Val	Asn	Leu	His	Leu	Leu
			195				200					205			
Leu	Leu	Arg	Asp	Ala	Ser	Ile	Phe	Gly	Glu	Glu	Trp	Gly	Phe	Thr	Pro
			210			215					220				
Gly	Glu	Ile	Ser	Arg	Phe	Tyr	Asn	Arg	Gln	Val	Gln	Leu	Thr	Ala	Glu
225					230					235					240
Tyr	Ser	Asp	Tyr	Cys	Val	Lys	Trp	Tyr	Lys	Ile	Gly	Leu	Asp	Lys	Leu
				245					250					255	
Lys	Gly	Thr	Thr	Ser	Lys	Ser	Trp	Leu	Asn	Tyr	His	Gln	Phe	Arg	Arg
			260					265					270		
Glu	Met	Thr	Leu	Leu	Val	Leu	Asp	Leu	Val	Ala	Leu	Phe	Pro	Asn	Tyr
			275				280					285			
Asp	Thr	His	Met	Tyr	Pro	Ile	Glu	Thr	Thr	Ala	Gln	Leu	Thr	Arg	Asp
			290			295					300				
Val	Tyr	Thr	Asp	Pro	Ile	Ala	Phe	Asn	Ile	Val	Thr	Ser	Thr	Gly	Phe
305					310					315					320
Cys	Asn	Pro	Trp	Ser	Thr	His	Ser	Gly	Ile	Leu	Phe	Tyr	Glu	Val	Glu
				325					330					335	
Asn	Asn	Val	Ile	Arg	Pro	Pro	His	Leu	Phe	Asp	Ile	Leu	Ser	Ser	Val
			340					345					350		
Glu	Ile	Asn	Thr	Ser	Arg	Gly	Gly	Ile	Thr	Leu	Asn	Asn	Asp	Ala	Tyr
			355				360					365			
Ile	Asn	Tyr	Trp	Ser	Gly	His	Thr	Leu	Lys	Tyr	Arg	Arg	Thr	Ala	Asp
			370			375					380				
Ser	Thr	Val	Thr	Tyr	Thr	Ala	Asn	Tyr	Gly	Arg	Ile	Thr	Ser	Glu	Lys
385					390					395					400
Asn	Ser	Phe	Ala	Leu	Glu	Asp	Arg	Asp	Ile	Phe	Glu	Ile	Asn	Ser	Thr
			405						410					415	
Val	Ala	Asn	Leu	Ala	Asn	Tyr	Tyr	Gln	Lys	Ala	Tyr	Gly	Val	Pro	Gly
			420					425					430		
Ser	Trp	Phe	His	Met	Val	Lys	Arg	Gly	Thr	Ser	Ser	Thr	Thr	Ala	Tyr
			435				440					445			
Leu	Tyr	Ser	Lys	Thr	His	Thr	Ala	Leu	Gln	Gly	Cys	Thr	Gln	Val	Tyr
			450			455					460				
Glu	Ser	Ser	Asp	Glu	Ile	Pro	Leu	Asp	Arg	Thr	Val	Pro	Val	Ala	Glu
465					470					475					480
Ser	Tyr	Ser	His	Arg	Leu	Ser	His	Ile	Thr	Ser	His	Ser	Phe	Ser	Lys
			485						490					495	
Asn	Gly	Ser	Ala	Tyr	Tyr	Gly	Ser	Phe	Pro	Val	Phe	Val	Trp	Thr	His
			500					505					510		
Thr	Ser	Ala	Asp	Leu	Asn	Asn	Thr	Ile	Tyr	Ser	Asp	Lys	Ile	Thr	Gln
			515				520					525			
Ile	Pro	Ala													

625					630					635				640
Ser	Met	Gly	Asp	Phe	Ser	Ser	Gly	Gln	Glu	Val	Tyr	Ile	Asp	Arg
				645					650				655	
Glu	Phe	Ile	Pro	Val	Asp	Glu	Thr	Tyr	Glu	Ala	Glu	Gln	Asp	Leu
			660					665					670	
Ala	Ala	Lys	Lys	Ala	Val	Asn	Ala	Leu	Phe	Thr	Asn	Thr	Lys	Asp
		675					680				685			Gly
Leu	Arg	Pro	Gly	Val	Thr	Asp	Tyr	Glu	Val	Asn	Gln	Ala	Ala	Asn
	690					695				700				Leu
Val	Glu	Cys	Leu	Ser	Asp	Asp	Leu	Tyr	Pro	Asn	Glu	Lys	Arg	Leu
705					710					715				Leu
Phe	Asp	Ala	Val	Arg	Glu	Ala	Lys	Arg	Leu	Ser	Gly	Ala	Arg	Asn
			725						730					Leu
Leu	Gln	Asp	Pro	Asp	Phe	Gln	Glu	Ile	Asn	Gly	Glu	Asn	Gly	Trp
		740					745					750		Ala
Ala	Ser	Thr	Gly	Ile	Glu	Ile	Val	Glu	Gly	Asp	Ala	Val	Phe	Lys
	755					760					765			Gly
Arg	Tyr	Leu	Arg	Leu	Pro	Gly	Ala	Arg	Glu	Ile	Asp	Thr	Glu	Thr
	770					775					780			Tyr
Pro	Thr	Tyr	Leu	Tyr	Gln	Lys	Val	Glu	Glu	Gly	Val	Leu	Lys	Pro
785					790					795				Tyr
Thr	Arg	Tyr	Arg	Leu	Arg	Gly	Phe	Val	Gly	Ser	Ser	Gln	Gly	Leu
			805						810					Glu
Ile	Tyr	Thr	Ile	Arg	His	Gln	Thr	Asn	Arg	Ile	Val	Lys	Asn	Val
		820						825					830	Pro
Asp	Asp	Leu	Leu	Pro	Asp	Val	Ser	Pro	Val	Asn	Ser	Asp	Gly	Ser
	835						840					845		Ile
Asn	Arg	Cys	Ser	Glu	Gln	Lys	Tyr	Val	Asn	Ser	Arg	Leu	Glu	Gly
	850					855					860			Glu
Asn	Arg	Ser	Gly	Asp	Ala	His	Glu	Phe	Ser	Leu	Pro	Ile	Asp	Ile
865					870					875				Gly
Glu	Leu	Asp	Tyr	Asn	Glu	Asn	Ala	Gly	Ile	Trp	Val	Gly	Phe	Lys
			885						890					Ile
Thr	Asp	Pro	Glu	Gly	Tyr	Ala	Thr	Leu	Gly	Asn	Leu	Glu	Leu	Val
		900						905				910		Glu
Glu	Gly	Pro	Leu	Ser	Gly	Asp	Ala	Leu	Glu	Arg	Leu	Gln	Arg	Glu
	915						920					925		Glu
Gln	Gln	Trp	Lys	Ile	Gln	Met	Thr	Arg	Arg	Arg	Glu	Glu	Thr	Asp
	930					935					940			Arg
Arg	Tyr	Met	Ala	Ser	Lys	Gln	Ala	Val	Asp	Arg	Leu	Tyr	Ala	Asp
945					950					955				Tyr
Gln	Asp	Gln	Gln	Leu	Asn	Pro	Asp	Val	Glu	Ile	Thr	Asp	Leu	Thr
			965						970					Ala
Ala	Gln	Asp	Leu	Ile	Gln	Ser	Ile	Pro	Tyr	Val	Tyr	Asn	Glu	Met
		980						985				990		Phe
Pro	Glu	Ile	Pro	Gly	Met	Asn	Tyr	Thr	Lys	Phe	Thr	Glu	Leu	Thr
	995						1000					1005		Asp
Arg	Leu	Gln	Gln	Ala	Trp	Asn	Leu	Tyr	Asp	Gln	Arg	Asn	Ala	Ile
	1010					1015					1020			Pro
Asn	Gly	Asp	Phe	Arg	Asn	Gly	Leu	Ser	Asn	Trp	Asn	Ala	Thr	Pro
1025					1030					1035				Gly
Val	Glu	Val	Gln	Gln	Ile	Asn	His	Thr	Ser	Val	Leu	Val	Ile	Pro
			1045						1050				1055	Asn
Trp	Asp	Glu	Gln	Val	Ser	Gln	Gln	Phe	Thr	Val	Gln	Pro	Asn	Gln
		1060						1065				1070		Arg
Tyr	Val	Leu	Arg	Val	Thr	Ala	Arg	Lys	Glu	Gly	Val	Gly	Asn	Gly
		1075					1080					1085		Tyr

Val	Ser	Ile	Arg	Asp	Gly	Gly	Asn	Gln	Ser	Glu	Thr	Leu	Thr	Phe	Ser
1090						1095					1100				
Ala	Ser	Asp	Tyr	Asp	Thr	Asn	Gly	Val	Tyr	Asn	Asp	Gln	Thr	Gly	Tyr
1105					1110					1115				1120	
Ile	Thr	Lys	Thr	Val	Thr	Phe	Ile	Pro	Tyr	Thr	Asp	Gln	Met	Trp	Ile
			1125					1130						1135	
Glu	Ile	Ser	Glu	Thr	Glu	Gly	Thr	Phe	Tyr	Ile	Glu	Ser	Val	Glu	Leu
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Ile	Val	Asp	Val	Glu											
		1155													

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<212> PRT

<213> *Bacillus thuringiensis*

<400> 19

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		20					25					30			
Asn	Lys	Pro	Asn	Gln	Pro	Leu	Lys	Asn	Thr	Asn	Tyr	Lys	Asp	Trp	Leu
	35					40					45				
Asn	Val	Cys	Gln	Asp	Asn	Gln	Gln	Tyr	Gly	Asn	Asn	Ala	Gly	Asn	Phe
50				55				60							
Ala	Ser	Ser	Glu	Thr	Ile	Val	Gly	Val	Ser	Ala	Gly	Ile	Ile	Val	Val
65				70				75						80	
Gly	Thr	Met	Leu	Gly	Ala	Phe	Ala	Ala	Pro	Val	Leu	Ala	Ala	Gly	Ile
			85					90						95	
Ile	Ser	Phe	Gly	Thr	Leu	Leu	Pro	Ile	Phe	Trp	Gln	Gly	Ser	Asp	Pro
		100					105						110		
Ala	Asn	Val	Trp	Gln	Asp	Leu	Leu	Asn	Ile	Gly	Gly	Arg	Pro	Ile	Gln
		115				120						125			
Glu	Ile	Asp	Lys	Asn	Ile	Ile	Asn	Val	Leu	Thr	Ser	Ile	Val	Thr	Pro
130						135					140				
Ile	Lys	Asn	Gln	Leu	Asp	Lys	Tyr	Gln	Glu	Phe	Phe	Asp	Lys	Trp	Glu
145				150						155				160	
Pro	Ala	Arg	Thr	His	Ala	Asn	Ala	Lys	Ala	Val	His	Asp	Leu	Phe	Thr
			165					170						175	
Thr	Leu	Glu	Pro	Ile	Ile	Asp	Lys	Asp	Leu	Asp	Met	Leu	Lys	Asn	Asn
		180						185					190		
Ala	Ser	Tyr	Arg	Ile	Pro	Thr	Leu	Pro	Ala	Tyr	Ala	Gln	Ile	Ala	Thr
		195				200						205			
Trp	His	Leu	Asn	Leu	Leu	Lys	His	Ala	Ala	Thr	Tyr	Tyr	Asn	Ile	Trp
210						215						220			
Leu	Gln	Asn	Gln	Gly	Ile	Asn	Pro	Ser	Thr	Phe	Asn	Ser	Ser	Asn	Tyr
225				230						235				240	
Tyr	Gln	Gly	Tyr	Leu	Lys	Arg	Lys	Ile	Gln	Glu	Tyr	Thr	Asp	Tyr	Cys
			245						250					255	
Ile	Gln	Thr	Tyr	Asn	Ala	Gly	Leu	Thr	Met	Ile	Arg	Thr	Asn	Thr	Asn
		260						265					270		
Ala	Thr	Trp	Asn	Met	Tyr	Asn	Thr	Tyr	Arg	Leu	Glu	Met	Thr	Leu	Thr
		275					280						285		
Val	Leu	Asp	Leu	Ile	Ala	Ile	Phe	Pro	Asn	Tyr	Asp	Pro	Glu	Lys	Tyr
290						295					300				
Pro	Ile	Gly	Val	Lys	Ser	Glu	Leu	Ile	Arg	Glu	Val	Tyr	Thr	Asn	Val

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Asn Ser Asp Thr Phe Arg Thr Ile Thr Glu Leu Glu Asn Gly Leu Thr						
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Arg Asn Pro Thr Leu Phe Thr Trp Ile Asn Gln Gly Arg Phe Tyr Thr						
	340			345		350
Arg Asn Ser Arg Asp Ile Leu Asp Pro Tyr Asp Ile Phe Ser Phe Thr						
	355			360		365
Gly Asn Gln Met Ala Phe Thr His Thr Asn Asp Asp Arg Asn Ile Ile						
	370			375		380
Trp Gly Ala Val His Gly Asn Ile Ile Ser Gln Asp Thr Ser Lys Val						
385		390			395	400
Phe Pro Phe Tyr Arg Asn Lys Pro Ile Asp Lys Val Glu Ile Val Arg						
	405			410		415
His Arg Glu Tyr Ser Asp Ile Ile Tyr Glu Met Ile Phe Phe Ser Asn						
	420			425		430
Ser Ser Glu Val Phe Arg Tyr Ser Ser Asn Ser Thr Ile Glu Asn Asn						
	435			440		445
Tyr Lys Arg Thr Asp Ser Tyr Met Ile Pro Lys Gln Thr Trp Lys Asn						
	450			455		460
Glu Glu Tyr Gly His Thr Leu Ser Tyr Ile Lys Thr Asp Asn Tyr Ile						
465		470			475	480
Phe Ser Val Val Arg Glu Arg Arg Arg Val Ala Phe Ser Trp Thr His						
	485			490		495
Thr Ser Val Asp Phe Gln Asn Thr Ile Asp Leu Asp Asn Ile Thr Gln						
	500			505		510
Ile His Ala Leu Lys Ala Leu Lys Val Ser Ser Asp Ser Lys Ile Val						
	515			520		525
Lys Gly Pro Gly His Thr Gly Gly Asp Leu Val Ile Leu Lys Asp Ser						
	530			535		540
Met Asp Phe Arg Val Arg Phe Leu Lys Asn Val Ser Arg Gln Tyr Gln						
545		550			555	560
Val Arg Ile Arg Tyr Ala Thr Asn Ala Pro Lys Thr Thr Val Phe Leu						
	565			570		575
Thr Gly Ile Asp Thr Ile Ser Val Glu Leu Pro Ser Thr Thr Ser Arg						
	580			585		590
Gln Asn Pro Asn Ala Thr Asp Leu Thr Tyr Ala Asp Phe Gly Tyr Val						
	595			600		605
Thr Phe Pro Arg Thr Val Pro Asn Lys Thr Phe Glu Gly Glu Asp Thr						
	610			615		620
Leu Leu Met Thr Leu Tyr Gly Thr Pro Asn His Ser Tyr Asn Ile Tyr						
625		630			635	640
Ile Asp Lys Ile Glu Phe Ile Pro Ile Thr Gln Ser Val Leu Asp Tyr						
	645			650		655
Thr Glu Lys Gln Asn Ile Glu Lys Thr Gln Lys Ile Val Asn Asp Leu						
	660			665		670
Phe Val Asn						
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 <212> PRT
 <213> Bacillus thuringiensis

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RTA01/2150211v1

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Asn	Gly	Glu	Thr	Phe	Gly	Gln	Lys	Arg	His	Gly	Tyr	Ser	Phe	Ala	Phe
				485					490					495	
Thr	His	Ser	Ser	Val	Asp	Pro	Asn	Asn	Thr	Ile	Ala	Ala	Asn	Lys	Ile
			500					505					510		
Thr	Gln	Ile	Pro	Val	Val	Lys	Ala	Ser	Ser	Ile	Asn	Gly	Ser	Ile	Ser
		515					520					525			
Ile	Glu	Lys	Gly	Pro	Gly	Phe	Thr	Gly	Gly	Asp	Leu	Val	Lys	Met	Arg
	530					535					540				
Ala	Asp	Ser	Gly	Leu	Thr	Met	Arg	Phe	Lys	Ala	Glu	Leu	Leu	Asp	Lys
545					550					555					560
Lys	Tyr	Arg	Val	Arg	Ile	Arg	Tyr	Lys	Cys	Asn	Tyr	Ser	Ser	Lys	Leu
				565					570					575	
Ile	Leu	Arg	Lys	Trp	Lys	Gly	Glu	Gly	Tyr	Ile	Gln	Gln	Gln	Ile	His
			580					585					590		
Asn	Ile	Ser	Pro	Thr	Tyr	Gly	Ala	Phe	Ser	Tyr	Leu	Glu	Ser	Phe	Thr
		595					600					605			
Ile	Thr	Thr	Thr	Glu	Asn	Ile	Phe	Asp	Leu	Thr	Met	Glu	Val	Thr	Tyr
	610					615					620				
Pro	Tyr	Gly	Arg	Gln	Phe	Val	Glu	Asp	Ile	Pro	Ser	Leu	Ile	Leu	Asp
625					630					635					640
Lys	Ile	Glu	Phe	Leu	Pro	Thr	Asn								
				645											

<210> 21

<211> 682

<212> PRT

<213> Bacillus thuringiensis

<400> 21

Met	Asn	Ser	Tyr	Gln	Asn	Lys	Asn	Glu	Tyr	Glu	Ile	Leu	Asp	Ala	Lys
1				5					10					15	
Arg	Asn	Thr	Cys	His	Met	Ser	Asn	Cys	Tyr	Pro	Lys	Tyr	Pro	Leu	Ala
			20					25					30		
Asn	Asp	Pro	Gln	Met	Tyr	Leu	Arg	Asn	Thr	His	Tyr	Lys	Asp	Trp	Ile
		35					40					45			
Asn	Met	Cys	Glu	Glu	Ala	Ser	Tyr	Ala	Ser	Ser	Gly	Pro	Ser	Gln	Leu
	50					55					60				
Phe	Lys	Val	Gly	Gly	Ser	Ile	Val	Ala	Lys	Ile	Leu	Gly	Met	Ile	Pro
65					70					75					80
Glu	Val	Gly	Pro	Leu	Leu	Ser	Trp	Met	Val	Ser	Leu	Phe	Trp	Pro	Thr
				85					90					95	
Ile	Glu	Glu	Lys	Asn	Thr	Val	Trp	Glu	Asp	Met	Ile	Lys	Tyr	Val	Ala
			100					105					110		
Asn	Leu	Leu	Lys	Gln	Glu	Leu	Thr	Asn	Asp	Thr	Leu	Asn	Arg	Ala	Thr
		115					120					125			
Ser	Asn	Leu	Ser	Gly	Leu	Asn	Glu	Ser	Leu	Asn	Ile	Tyr	Asn	Arg	Ala
	130					135					140				
Leu	Ala	Ala	Trp	Lys	Gln	Asn	Lys	Asn	Asn	Phe	Ala	Ser	Gly	Glu	Leu
145					150					155					160
Ile	Arg	Ser	Tyr	Ile	Asn	Asp	Leu	His	Ile	Leu	Phe	Thr	Arg	Asp	Ile
				165					170					175	
Gln	Ser	Asp	Phe	Ser	Leu	Gly	Gly	Tyr	Glu	Thr	Val	Leu	Leu	Pro	Ser
			180					185					190		
Tyr	Ala	Ser	Ala	Ala	Asn	Leu	His	Leu	Leu	Leu	Leu	Arg	Asp	Val	Ala
		195					200					205			

Ile	Tyr	Gly	Lys	Glu	Leu	Gly	Tyr	Pro	Ser	Thr	Asp	Val	Glu	Phe	Tyr	210	215	220
Tyr	Asn	Glu	Gln	Lys	Tyr	Tyr	Thr	Glu	Lys	Tyr	Ser	Asn	Tyr	Cys	Val	225	230	235
Asn	Thr	Tyr	Lys	Ser	Gly	Leu	Glu	Ser	Lys	Lys	Gln	Ile	Gly	Trp	Ser	245	250	255
Asp	Phe	Asn	Arg	Tyr	Arg	Arg	Glu	Met	Thr	Leu	Ser	Val	Leu	Asp	Ile	260	265	270
Val	Ala	Leu	Phe	Pro	Leu	Tyr	Asp	Thr	Gly	Leu	Tyr	Pro	Ser	Lys	Asp	275	280	285
Gly	Lys	Ile	His	Val	Lys	Ala	Glu	Leu	Thr	Arg	Glu	Ile	Tyr	Ser	Asp	290	295	300
Val	Ile	Asn	Asp	His	Val	Tyr	Gly	Leu	Met	Val	Pro	Tyr	Ile	Ser	Phe	305	310	315
Glu	His	Ala	Glu	Ser	Leu	Tyr	Thr	Arg	Arg	Pro	His	Ala	Phe	Thr	Trp	325	330	335
Leu	Lys	Gly	Phe	Arg	Phe	Val	Thr	Asn	Ser	Ile	Asn	Ser	Trp	Thr	Phe	340	345	350
Leu	Ser	Gly	Gly	Glu	Asn	Arg	Tyr	Phe	Leu	Thr	His	Gly	Glu	Gly	Thr	355	360	365
Ile	Tyr	Asn	Gly	Pro	Phe	Leu	Gly	Gln	Asp	Thr	Glu	Tyr	Gly	Gly	Thr	370	375	380
Ser	Ser	Tyr	Ile	Asp	Ile	Ser	Asn	Asn	Ser	Ser	Ile	Tyr	Asn	Leu	Trp	385	390	395
Thr	Lys	Asn	Tyr	Glu	Trp	Ile	Tyr	Pro	Trp	Thr	Asp	Pro	Val	Asn	Ile	405	410	415
Thr	Lys	Ile	Asn	Phe	Ser	Ile	Thr	Asp	Asn	Ser	Asn	Ser	Ser	Glu	Ser	420	425	430
Ile	Tyr	Gly	Ala	Glu	Arg	Met	Asn	Lys	Pro	Thr	Val	Arg	Thr	Asp	Phe	435	440	445
Asn	Phe	Leu	Leu	Asn	Arg	Ala	Gly	Asn	Gly	Pro	Thr	Thr	Tyr	Asn	Asp	450	455	460
Tyr	Asn	His	Ile	Leu	Ser	Tyr	Met	Leu	Ile	Asn	Gly	Glu	Thr	Phe	Gly	465	470	475
Gln	Lys	Arg	His	Gly	Tyr	Ser	Phe	Ala	Phe	Thr	His	Ser	Ser	Val	Asp	485	490	495
Arg	Tyr	Asn	Thr	Ile	Val	Pro	Asp	Lys	Ile	Val	Gln	Ile	Pro	Ala	Val	500	505	510
Lys	Thr	Asn	Leu	Val	Gly	Ala	Asn	Ile	Ile	Lys	Gly	Pro	Gly	His	Thr	515	520	525
Gly	Gly	Asp	Leu	Leu	Lys	Leu	Glu	Tyr	Glu	Arg	Phe	Leu	Ser	Leu	Arg	530	535	540
Ile	Lys	Leu	Ile	Ala	Ser	Met	Thr	Phe	Arg	Ile	Arg	Ile	Arg	Tyr	Ala	545	550	555
Ser	Asn	Ile	Ser	Gly	Gln	Met	Met	Ile	Asn	Ile	Gly	Tyr	Gln	Asn	Pro	565	570	575
Thr	Tyr	Phe	Asn	Ile	Ile	Pro	Thr	Thr	Ser	Arg	Asp	Tyr	Thr	Glu	Leu	580	585	590
Lys	Phe	Glu	Asp	Phe	Gln	Leu	Val	Asp	Thr	Ser	Tyr	Ile	Tyr	Ser	Gly	595	600	605
Gly	Pro	Ser	Ile	Ser	Ser	Asn	Thr	Leu	Trp	Leu	Asp	Asn	Phe	Ser	Asn	610	615	620
Gly	Pro	Val	Ile	Ile	Asp	Lys	Ile	Glu	Phe	Ile	Pro	Leu	Gly	Ile	Thr	625	630	635
Leu	Asn	Gln	Ala	Gln	Gly	Tyr	Asp	Thr	Tyr	Asp	Gln	Asn	Ala	Asn	Gly	645	650	655
Met	Tyr	His	Gln	Asn	Tyr	Ser	Asn	Ser	Gly	Tyr	Asn	Tyr	Asn	Gln	Glu			

	660		665		670
Tyr	Asn	Thr	Tyr	Tyr	Gln
	675		680	Asn	Asn

<210> 22
 <211> 674
 <212> PRT
 <213> *Bacillus thuringiensis*

<400> 22

Met	Asn	Gln	Tyr	Gln	Asn	Lys	Asn	Glu	Tyr	Glu	Ile	Leu	Glu	Ser	Ser
1				5				10					15		
Gln	Asn	Asn	Met	Asn	Met	Pro	Asn	Arg	Tyr	Pro	Phe	Ala	Asp	Asp	Pro
		20					25					30			
Asn	Ala	Val	Met	Lys	Asn	Gly	Asn	Tyr	Lys	Asp	Trp	Val	Asn	Glu	Cys
	35					40					45				
Glu	Gly	Ser	Asn	Ile	Ser	Pro	Ser	Pro	Ala	Ala	Ala	Ile	Thr	Ser	Lys
	50					55				60					
Ile	Val	Ser	Ile	Val	Leu	Lys	Thr	Leu	Ala	Lys	Ala	Val	Ala	Ser	Ser
65					70					75					80
Leu	Ala	Asp	Ser	Ile	Lys	Ser	Ser	Leu	Gly	Ile	Ser	Lys	Thr	Ile	Thr
				85				90						95	
Glu	Asn	Asn	Val	Ser	Gln	Val	Ser	Met	Val	Gln	Val	His	Gln	Ile	Ile
			100					105					110		
Asn	Arg	Arg	Ile	Gln	Glu	Thr	Ile	Leu	Asp	Leu	Gly	Glu	Ser	Ser	Leu
		115					120					125			
Asn	Gly	Leu	Val	Ala	Ile	Tyr	Asn	Arg	Asp	Tyr	Leu	Gly	Ala	Leu	Glu
	130					135					140				
Ala	Trp	Asn	Asn	Asn	Lys	Ser	Asn	Ile	Asn	Tyr	Gln	Thr	Asn	Val	Ala
145					150					155					160
Glu	Ala	Phe	Lys	Thr	Val	Glu	Arg	Glu	Phe	Phe	Thr	Lys	Leu	Lys	Gly
				165					170					175	
Ile	Tyr	Arg	Thr	Ser	Ser	Ser	Gln	Ile	Thr	Leu	Leu	Pro	Thr	Phe	Thr
			180					185						190	
Gln	Ala	Ala	Asn	Leu	His	Leu	Ser	Met	Leu	Arg	Asp	Ala	Val	Met	Tyr
		195					200					205			
Gln	Glu	Gly	Trp	Asn	Leu	Gln	Ser	His	Ile	Asn	Tyr	Ser	Lys	Glu	Leu
	210					215					220				
Asp	Asp	Ala	Leu	Glu	Asp	Tyr	Thr	Asn	Tyr	Cys	Val	Glu	Val	Tyr	Thr
225					230					235					240
Lys	Gly	Leu	Asn	Ala	Leu	Arg	Gly	Ser	Thr	Ala	Ile	Asp	Trp	Leu	Glu
				245					250					255	
Phe	Asn	Ser	Phe	Arg	Arg	Asp	Met	Thr	Leu	Met	Val	Leu	Asp	Leu	Val
			260					265					270		
Ala	Ile	Phe	Pro	Asn	Tyr	Asn	Pro	Val	Arg	Tyr	Pro	Leu	Ser	Thr	Lys
		275					280						285		
Ile	Ser	Leu	Ser	Arg	Lys	Ile	Tyr	Thr	Asp	Pro	Val	Gly	Arg	Thr	Asp
	290					295					300				
Ser	Pro	Ser	Phe	Gly	Asp	Trp	Thr	Asn	Thr	Gly	Arg	Thr	Leu	Ala	Asn
305					310					315					320
Phe	Asn	Asp	Leu	Glu	Arg	Glu	Val	Thr	Asp	Ser	Pro	Ser	Leu	Val	Lys
				325					330					335	
Trp	Leu	Gly	Asp	Met	Thr	Ile	Tyr	Thr	Gly	Ala	Ile	Asp	Ser	Tyr	Arg
			340					345					350		
Pro	Thr	Ser	Pro	Gly	Asp	Arg	Ile	Gly	Val	Trp	Tyr	Gly	Asn	Ile	Asn
		355					360						365		

Ala Phe Tyr His Thr Gly Arg Thr Asp Val Val Met Phe Arg Gln Thr
 370 375 380
 Gly Asp Thr Ala Tyr Glu Asp Pro Ser Thr Phe Ile Ser Asn Ile Leu
 385 390 395 400
 Tyr Asp Asp Ile Tyr Lys Leu Asp Leu Arg Ala Ala Ala Val Ser Thr
 405 410 415
 Ile Gln Gly Ala Met Asp Thr Thr Phe Gly Val Ser Ser Ser Arg Phe
 420 425 430
 Phe Asp Ile Arg Gly Arg Asn Gln Leu Tyr Gln Ser Asn Lys Pro Tyr
 435 440 445
 Pro Ser Leu Pro Ile Thr Ile Thr Phe Pro Gly Glu Glu Ser Ser Glu
 450 455 460
 Gly Asn Ala Asn Asp Tyr Ser His Leu Leu Cys Asp Val Lys Ile Leu
 465 470 475 480
 Gln Glu Asp Ser Ser Asn Ile Cys Glu Gly Arg Ser Ser Leu Leu Ser
 485 490 495
 His Ala Trp Thr His Ala Ser Leu Asp Arg Asn Asn Thr Ile Leu Pro
 500 505 510
 Asp Glu Ile Thr Gln Ile Pro Ala Val Thr Ala Tyr Glu Leu Arg Gly
 515 520 525
 Asn Ser Ser Val Val Ala Gly Pro Gly Ser Thr Gly Gly Asp Leu Val
 530 535 540
 Lys Met Ser Tyr His Ser Val Trp Ser Phe Lys Val Tyr Cys Ser Glu
 545 550 555 560
 Leu Lys Asn Tyr Arg Val Arg Ile Arg Tyr Ala Ser His Gly Asn Cys
 565 570 575
 Gln Phe Leu Met Lys Arg Trp Pro Ser Thr Gly Val Ala Pro Arg Gln
 580 585 590
 Trp Ala Arg His Asn Val Gln Gly Thr Phe Ser Asn Ser Met Arg Tyr
 595 600 605
 Glu Ala Phe Lys Tyr Leu Asp Ile Phe Thr Ile Thr Pro Glu Glu Asn
 610 615 620
 Asn Phe Ala Phe Thr Ile Asp Leu Glu Ser Gly Gly Asp Leu Phe Ile
 625 630 635 640
 Asp Lys Ile Glu Phe Ile Pro Val Ser Gly Ser Ala Phe Glu Tyr Glu
 645 650 655
 Gly Lys Gln Asn Ile Glu Lys Thr Gln Lys Ala Val Asn Asp Leu Phe
 660 665 670
 Ile Asn

<210> 23
 <211> 675
 <212> PRT
 <213> *Bacillus thuringiensis*

<400> 23
 Met Asn Pro Tyr Gln Asn Lys Ser Glu Cys Glu Ile Leu Asn Ala Pro
 1 5 10 15
 Leu Asn Asn Ile Asn Met Pro Asn Arg Tyr Pro Phe Ala Asn Asp Pro
 20 25 30
 Asn Ala Val Met Lys Asn Gly Asn Tyr Lys Asp Trp Leu Asn Glu Cys
 35 40 45
 Asp Gly Ile Thr Pro Ser Ile Phe Gly Thr Leu Gly Val Leu Ala Ser
 50 55 60
 Ile Val Ile Ser Thr Ile Asn Leu Ala Thr Ser Pro Ser Ile Gly Asp

65					70					75					80
Ala	Phe	Ala	Leu	Val	Ser	Ser	Ile	Gly	Glu	Tyr	Trp	Pro	Glu	Thr	Lys
				85					90					95	
Thr	Ser	Phe	Pro	Leu	Ser	Val	Ala	Asp	Val	Asn	Arg	Leu	Ile	Arg	Glu
			100					105					110		
Ala	Leu	Asp	Gln	Asn	Ala	Ile	Asn	Arg	Ala	Thr	Gly	Lys	Phe	Asn	Gly
		115					120					125			
Leu	Met	Asp	Thr	Tyr	Asn	Thr	Val	Tyr	Leu	Lys	Asn	Leu	Gln	Asp	Trp
	130					135					140				
Tyr	Asp	Thr	Arg	Ile	Pro	Ala	Asn	Pro	Gln	Gly	Asp	Ser	Gln	Leu	Arg
145					150					155					160
Glu	Ala	Ala	Arg	Arg	Ser	Leu	Glu	Glu	Ile	Glu	Arg	Asp	Phe	Arg	Lys
			165					170						175	
Ala	Leu	Ala	Gly	Glu	Phe	Ala	Glu	Ala	Gly	Ser	Gln	Ile	Val	Leu	Leu
		180					185					190			
Pro	Ile	Tyr	Ala	Gln	Ala	Ala	Asn	Ile	His	Leu	Leu	Ile	Leu	Lys	Asp
	195						200					205			
Ala	Met	Gln	Phe	Arg	Thr	Asp	Leu	Gly	Leu	Ile	Arg	Pro	Val	Gly	Val
	210				215						220				
Pro	Ile	Thr	Thr	Ser	Ala	Glu	Asp	Pro	Phe	Glu	Ser	Glu	Phe	Leu	Leu
225				230					235						240
Arg	Ile	Lys	Lys	Tyr	Thr	Asp	His	Cys	Ile	Ser	Tyr	Tyr	Asp	Asp	Gly
			245				250						255		
Leu	Ala	Lys	Ile	Arg	Ser	Arg	Gly	Ser	Asp	Gly	Glu	Thr	Trp	Trp	Glu
		260					265					270			
Phe	Asn	Lys	Phe	Arg	Arg	Glu	Met	Thr	Leu	Thr	Val	Leu	Asp	Leu	Val
	275					280					285				
Ala	Leu	Tyr	Pro	Thr	His	Asn	Ile	Lys	Leu	Tyr	Pro	Ile	Pro	Thr	Gln
	290				295						300				
Thr	Glu	Leu	Ser	Arg	Val	Val	Tyr	Thr	Asp	Pro	Val	Gly	Cys	Phe	Gly
305				310					315						320
Asn	Arg	Lys	Ser	Asp	Ile	Phe	Ser	Arg	Leu	Asn	Phe	Asp	Tyr	Leu	Glu
			325					330					335		
Asn	Arg	Leu	Thr	Arg	Pro	Arg	Glu	Pro	Phe	Asn	Tyr	Leu	Asn	Ser	Val
		340					345					350			
Gln	Leu	Phe	Ala	Ser	Thr	Val	Ser	Asn	Ser	Asn	Asn	Gly	Glu	Val	Leu
	355					360					365				
Arg	Gly	Asn	Leu	Asn	Lys	Ile	Met	Phe	Glu	Gly	Gly	Trp	Thr	Ala	Ser
	370				375						380				
Arg	Ser	Gly	Asp	Gly	Val	Thr	Thr	Gly	Thr	Pro	Phe	Ser	Thr	Met	Asp
385				390					395					400	
Trp	Ser	Tyr	Gly	Trp	Gly	Tyr	Pro	Arg	Lys	His	Tyr	Ala	Glu	Ile	Thr
			405				410						415		
Ser	Arg	Ser	Gln	Ala	Leu	Pro	Gly	Leu	Asn	Asn	Ser	Ile	His	Val	Ile
		420					425					430			
Val	Gly	Ile	Asp	Ser	Phe	Arg	Ala	Ile	Gly	Pro	Gly	Gly	Gln	Gly	Asp
	435					440					445				
His	Thr	Phe	Ser	Leu	Pro	Gly	Gly	Asp	Met	Tyr	Asp	Cys	Gly	Lys	Val
	450				455					460					
Gln	Ile	Asn	Pro	Leu	Glu	Asp	Tyr	Arg	Asn	Ser	Asp	His	Trp	Ile	Ser
465				470					475					480	
Asp	Met	Met	Thr	Ile	Asn	Gln	Ser	Val	Gln	Leu	Ala	Ser	Asn	Pro	Thr
			485				490						495		
Gln	Thr	Phe	Ala	Phe	Ser	Ala	Leu	Ser	Leu	Gly	Trp	His	His	Ser	Ser
		500					505					510			
Ala	Gly	Asn	Arg	Asn	Val	Tyr	Val	Tyr	Asp	Lys	Ile	Thr	Gln	Ile	Pro
	515						520					525			

Ala	Thr	Lys	Thr	Val	Arg	Glu	His	Pro	Met	Ile	Lys	Gly	Pro	Gly	Phe
530						535				540					
Thr	Gly	Gly	Asp	Leu	Ala	Asp	Leu	Ser	Ser	Asn	Ser	Asp	Ile	Leu	Gln
545					550					555					560
Tyr	Asp	Leu	Arg	Ser	Asp	Tyr	Asp	Asp	Arg	Leu	Thr	Glu	Asp	Val	Pro
				565					570					575	
Phe	Arg	Ile	Arg	Ile	Arg	Cys	Ala	Ser	Ile	Gly	Val	Ser	Thr	Ile	Ser
			580					585					590		
Val	Asp	Asn	Trp	Gly	Ser	Ser	Ser	Pro	Gln	Val	Thr	Val	Ala	Ser	Thr
	595						600					605			
Ala	Ala	Ser	Leu	Asp	Thr	Leu	Lys	Tyr	Glu	Ser	Phe	Gln	Tyr	Val	Ser
610						615					620				
Ile	Pro	Gly	Asn	Tyr	Tyr	Phe	Asp	Ser	Ala	Pro	Arg	Ile	Arg	Leu	Leu
625					630					635					640
Arg	Gln	Pro	Gly	Arg	Leu	Leu	Val	Asp	Arg	Ile	Glu	Ile	Ile	Pro	Val
				645					650					655	
Asn	Phe	Phe	Pro	Leu	Ser	Glu	Gln	Glu	Asn	Lys	Ser	Val	Asp	Ser	Leu
			660					665					670		
Phe	Ile	Asn													
		675													

<210> 24

<211> 659

<212> PRT

<213> *Bacillus thuringiensis*

<400> 24

Asn	Ser	Tyr	Glu	Asn	Lys	Asn	Glu	Tyr	Glu	Ile	Leu	Asn	Asp	Ser	Lys
1				5					10					15	
Lys	Ser	Asn	Met	Ser	Asn	Pro	Tyr	Leu	Arg	Tyr	Pro	Leu	Ala	Asn	Asp
			20					25				30			
Ser	Leu	Ala	Ser	Met	Gln	Asn	Thr	Asn	Tyr	Lys	Asp	Trp	Leu	Thr	Met
		35					40					45			
Cys	Asp	Arg	Thr	Asp	Thr	Asp	Val	Leu	Ser	Ser	Arg	Gly	Ala	Val	Ser
	50					55					60				
Thr	Gly	Val	Gly	Met	Leu	Ser	Thr	Ile	Leu	Ser	Leu	Phe	Gly	Ile	Pro
65					70					75					80
Leu	Ile	Gly	Glu	Gly	Ile	Asp	Leu	Leu	Leu	Gly	Ala	Ala	Asp	Phe	Leu
				85					90					95	
Trp	Pro	Glu	Ser	Asp	Gly	Gly	His	Gln	Tyr	Thr	Trp	Glu	Asp	Leu	Met
			100					105					110		
Asn	His	Ile	Glu	Glu	Leu	Met	Asp	Glu	Arg	Leu	Glu	Thr	Glu	Lys	Arg
		115					120					125			
Thr	Thr	Ala	Leu	Asp	Asp	Leu	Arg	Gly	Leu	Lys	Ala	Leu	Leu	Gly	Leu
	130					135					140				
Phe	Arg	Asp	Ala	Phe	Asp	Ser	Trp	Glu	Lys	Asn	Gln	Asn	Asp	Pro	Ile
145					150					155					160
Ala	Lys	Asn	Arg	Val	Gly	Gly	Tyr	Phe	Glu	Asp	Val	His	Thr	His	Phe
				165					170					175	
Val	Lys	Asp	Met	Ala	Ser	Ile	Phe	Ser	Ala	Thr	Asn	Tyr	Glu	Val	Leu
			180					185					190		
Leu	Leu	Pro	Val	Tyr	Ala	Gln	Ala	Ala	Asn	Leu	His	Leu	Leu	Leu	Leu
		195					200					205			
Arg	Glu	Gly	Val	Ile	Tyr	Gly	Ser	Arg	Trp	Gly	Ile	Ala	Pro	Ala	Ala
	210					215					220				
Asp	Phe	Tyr	His	Asp	Gln	Leu	Leu	Lys	Tyr	Thr	Ala	Ile	Tyr	Ala	Asn

225					230					235				240
His	Cys	Val	Thr	Trp	Tyr	Asn	Asn	Gly	Leu	Ala	Gln	Gln	Lys	Glu
				245					250					255
Phe	Ala	Lys	Ser	Pro	Asn	Trp	Asn	Arg	Phe	Asn	Ala	Tyr	Arg	Arg
			260					265					270	
Met	Thr	Ile	Thr	Val	Leu	Asp	Ile	Ile	Ala	Leu	Phe	Pro	Thr	Tyr
		275					280					285		
Ala	Arg	Leu	Tyr	Thr	Lys	Pro	Ile	Lys	Thr	Glu	Leu	Thr	Arg	Glu
	290					295				300				
Tyr	Ser	Asp	Val	Leu	Asn	Leu	Asp	Val	Tyr	Gly	Val	Gln	Gln	Thr
305					310					315				320
Leu	Asn	Lys	Asn	Glu	Ala	Ala	Phe	Thr	Arg	Ser	Pro	His	Leu	Val
			325					330					335	
Arg	Leu	Arg	Gly	Phe	Asp	Phe	Tyr	Thr	Arg	Thr	Lys	Tyr	Ala	Tyr
			340					345					350	
Arg	Tyr	Leu	Ala	Gly	His	Thr	Asn	Tyr	Phe	Ser	Phe	Thr	Gly	Asn
		355					360					365		
Thr	Ile	Tyr	Ser	Ser	Ser	Phe	Asn	Asn	Trp	Tyr	Asp	Thr	Asp	Met
	370					375					380			
Lys	Ser	Thr	Ile	Asn	Ile	Pro	Asp	Tyr	Ala	Asn	Ile	Tyr	Lys	Leu
385					390					395				400
Thr	Lys	Ser	Tyr	Thr	Asn	Ile	Ser	Pro	Tyr	Thr	Asp	Pro	Val	Gly
				405					410				415	
Ser	Gln	Met	Gln	Phe	Ser	Leu	Thr	Asn	Asn	Gln	Gln	Leu	Thr	Tyr
			420					425				430		
Gly	Thr	Ser	Ala	Pro	Lys	Tyr	Pro	Val	Arg	Glu	Thr	Phe	Phe	Glu
		435					440					445		
Pro	Pro	Thr	Asp	Glu	Lys	Pro	Leu	Thr	Tyr	Glu	Asn	Tyr	Ser	His
	450					455					460			
Leu	Ser	Tyr	Met	Thr	Ser	Ala	Gln	His	Phe	Gly	Asp	Lys	Lys	Ile
465					470					475				480
Tyr	Thr	Phe	Ala	Trp	Met	His	Glu	Ser	Val	Asp	Phe	Asp	Asn	Arg
			485					490					495	
Asp	Pro	Asp	Lys	Ile	Thr	Gln	Ile	Pro	Ala	Val	Lys	Gly	Asp	Tyr
			500					505					510	
Gln	Tyr	Gly	Tyr	Val	Lys	Gln	Gly	Pro	Gly	His	Thr	Gly	Gly	Asp
		515					520					525		
Val	Ser	Met	Ile	Arg	Thr	Asp	Arg	Leu	Gly	Ile	Asn	Val	Tyr	Phe
	530					535					540			
Gln	Pro	Leu	Asp	Tyr	Arg	Ile	Arg	Ile	Arg	Tyr	Ser	Thr	Ser	Ser
545					550					555				560
Gly	Tyr	Leu	Tyr	Ile	Tyr	Ser	Pro	Asn	Thr	Lys	Ile	Val	Tyr	Leu
				565					570					575
Pro	Thr	Thr	Leu	Val	Asp	Gly	Gln	Pro	Thr	Phe	Asp	Pro	Met	Asp
			580					585					590	
Ser	Ala	Phe	Arg	Val	Val	Glu	Val	Pro	Ala	Ser	Phe	Arg	Ala	Ser
		595					600					605		
Ala	Gly	Tyr	Thr	Asn	Phe	Thr	Ile	Glu	Ala	Gly	Phe	Gly	Pro	Val
	610					615					620			
Ile	Asp	Lys	Ile	Glu	Phe	Ile	Pro	Asp	Asn	Thr	Thr	Thr	Leu	Glu
625					630					635				640
Glu	Gly	Gly	Arg	Asp	Leu	Glu	Lys	Thr	Lys	Asn	Ala	Val	Asn	Asp
			645						650					655
Phe	Thr	Asn												

<210> 25
 <211> 666
 <212> PRT
 <213> *Bacillus thuringiensis*

<400> 25

Asn	Ser	Tyr	Glu	Asn	Lys	Asn	Glu	Tyr	Glu	Ile	Leu	Glu	Ser	Ser	Ser
1				5					10					15	
Asn	Asn	Thr	Asn	Met	Pro	Asn	Arg	Tyr	Pro	Phe	Ala	Asn	Asp	Arg	Asp
			20					25					30		
Met	Ser	Thr	Met	Ser	Phe	Asn	Asp	Cys	Gln	Gly	Ile	Ser	Trp	Asp	Glu
		35					40						45		
Ile	Trp	Glu	Ser	Ala	Glu	Thr	Ile	Thr	Ser	Ile	Gly	Ile	Asp	Leu	Ile
	50					55					60				
Glu	Phe	Leu	Met	Glu	Pro	Ser	Leu	Gly	Gly	Ile	Asn	Thr	Leu	Phe	Ser
65					70					75					80
Ile	Ile	Gly	Lys	Leu	Ile	Pro	Thr	Asn	His	Gln	Ser	Val	Ser	Ala	Leu
				85					90					95	
Ser	Ile	Cys	Asp	Leu	Leu	Ser	Ile	Ile	Arg	Lys	Glu	Val	Ala	Asp	Ser
			100					105					110		
Val	Leu	Ser	Asp	Ala	Ile	Cys	Arg	Phe	Leu	Asp	Gly	Lys	Leu	Lys	Asn
		115					120					125			
Tyr	Arg	Glu	Tyr	Tyr	Leu	Pro	Tyr	Leu	Glu	Ala	Trp	Leu	Lys	Asp	Gly
	130					135						140			
Lys	Pro	Leu	Gln	Lys	Thr	Asn	Asn	Ser	Asp	Ile	Gly	Gln	Leu	Val	Lys
145					150					155					160
Tyr	Phe	Glu	Leu	Ser	Glu	Arg	Asp	Phe	Asn	Glu	Ile	Leu	Gly	Gly	Ser
				165					170					175	
Leu	Ala	Arg	Asn	Asn	Ala	Gln	Ile	Leu	Leu	Leu	Pro	Tyr	Phe	Cys	Ala
			180					185					190		
Ser	Cys	Lys	Cys	Gln	Leu	Leu	Leu	Arg	Asp	Ala	Val	Gln	Tyr	Glu	
		195					200				205				
Glu	Gln	Trp	Phe	Pro	Phe	Leu	Ser	Ala	Glu	Asn	Val	Arg	Ser	Glu	Leu
	210					215					220				
Ile	Ser	Pro	Asn	Ser	Gly	Cys	Asp	Phe	Thr	Gly	Asp	Tyr	Tyr	Glu	Arg
225					230					235					240
Leu	Lys	Cys	Lys	Ile	Ala	Glu	Tyr	Thr	Asp	Tyr	Cys	Glu	Tyr	Trp	Tyr
				245					250					255	
Gln	Ala	Gly	Leu	Asn	Gln	Ile	Lys	Gln	Ala	Gly	Thr	Gly	Ala	Asp	Thr
			260					265					270		
Trp	Ala	Lys	Phe	Asn	Lys	Phe	Arg	Arg	Glu	Met	Thr	Leu	Thr	Val	Leu
		275					280					285			
Asp	Ile	Ile	Ala	Ile	Phe	Gln	Thr	Tyr	Asp	Phe	Lys	Lys	Tyr	Pro	Leu
	290					295					300				
Pro	Thr	His	Val	Glu	Leu	Thr	Arg	Glu	Ile	Tyr	Thr	Asp	Pro	Val	Gly
305					310					315					320
Tyr	Ser	Ser	Gly	Thr	Tyr	Ser	Trp	Leu	Lys	Tyr	Trp	Thr	Gly	Ala	Phe
			325						330					335	
Asn	Thr	Leu	Glu	Ala	Asn	Gly	Thr	Arg	Gly	Pro	Gly	Leu	Val	Thr	Trp
			340					345					350		
Leu	Arg	Ser	Ile	Gly	Ile	Tyr	Asn	Glu	Tyr	Val	Ser	Arg	Tyr	Phe	Ser
		355					360					365			
Gly	Trp	Val	Gly	Thr	Arg	His	Tyr	Glu	Asp	Tyr	Thr	Thr	Gly	Asn	Gly
	370					375					380				
Asn	Phe	Gln	Arg	Met	Ser	Gly	Thr	Thr	Ser	Asn	Asp	Leu	Arg	Asp	Ile
385					390					395					400
Ser	Phe	Pro	Asn	Ser	Asp	Ile	Phe	Lys	Ile	Glu	Ser	Lys	Ala	Ile	Met

				405					410					415	
Asn	Leu	Val	Gly	Glu	Ile	Asn	Ala	Arg	Pro	Glu	Tyr	Arg	Val	Ser	Arg
			420					425					430		
Ala	Glu	Phe	Ser	Glu	Ser	Thr	Ala	Phe	Ile	Tyr	Leu	Tyr	Asp	Ala	Gly
		435					440					445			
Asn	Ser	Gly	Leu	Ser	Ser	Met	Thr	Ile	Thr	Ser	Lys	Leu	Pro	Gly	Ile
	450					455					460				
Lys	Asn	Pro	Glu	Pro	Ser	Tyr	Arg	Asp	Tyr	Ser	His	Arg	Leu	Ser	Asn
465					470					475					480
Ala	Ala	Cys	Val	Gly	Ala	Gly	Asn	Ser	Arg	Ile	Asn	Val	Tyr	Gly	Trp
			485						490					495	
Thr	His	Thr	Ser	Met	Ser	Lys	Tyr	Asn	Leu	Ile	Tyr	Pro	Asp	Lys	Ile
			500					505					510		
Thr	Gln	Ile	Pro	Ala	Val	Lys	Ala	Phe	Asp	Ile	Ser	Asp	Thr	Gly	Pro
	515						520					525			
Gly	Gln	Val	Ile	Ala	Gly	Pro	Gly	His	Thr	Gly	Gly	Asn	Val	Val	Ser
	530					535					540				
Leu	Pro	Tyr	Tyr	Ser	Arg	Leu	Lys	Ile	Arg	Leu	Ile	Pro	Ala	Ser	Thr
545					550					555					560
Asn	Lys	Asn	Tyr	Leu	Val	Arg	Val	Arg	Tyr	Thr	Ser	Thr	Ser	Asn	Gly
			565					570						575	
Arg	Leu	Leu	Val	Glu	Arg	Trp	Ser	Pro	Ser	Ser	Ile	Ile	Asn	Ser	Tyr
			580					585					590		
Phe	Phe	Leu	Pro	Ser	Thr	Gly	Pro	Gly	Asp	Ser	Phe	Gly	Tyr	Val	Asp
		595					600					605			
Thr	Leu	Val	Thr	Thr	Phe	Asn	Gln	Pro	Gly	Val	Glu	Ile	Ile	Ile	Gln
	610					615					620				
Asn	Leu	Asp	Thr	Pro	Ile	Asn	Val	Asp	Lys	Val	Glu	Phe	Ile	Pro	Val
625					630					635					640
Asn	Ser	Thr	Ala	Leu	Glu	Tyr	Glu	Gly	Lys	Gln	Ser	Leu	Glu	Lys	Ala
			645					650						655	
Gln	Asp	Val	Val	Asn	Asp	Leu	Phe	Val	Lys						
		660						665							

<210> 26
 <211> 529
 <212> PRT
 <213> Bacillus thuringiensis

<400> 26
 Val Asn Phe Met Leu Thr Ser Gly Ala Lys Asn Met Leu Lys Leu Glu
 1 5 10 15
 Thr Thr Asp Tyr Glu Ile Asp Gln Met Ala Asn Ala Ile Glu Asn Met
 20 25 30
 Ser Gly Glu Gln Tyr Ser Gln Glu Lys Met Met Gln Trp His Asp Ile
 35 40 45
 Lys Tyr Ala Lys Gln Leu Ser Gln Ala Arg Asn Leu Leu Gln Asn Gly
 50 55 60
 Asp Phe Glu Asp Leu Phe Ser Gly Trp Thr Thr Ser Asn Gln Met Ser
 65 70 75 80
 Ile Gln Ala Asp Asn Ala Thr Phe Lys Gly Asn Tyr Leu His Met Ser
 85 90 95
 Gly Ala Arg Asp Ile Tyr Gly Thr Ile Phe Pro Thr Tyr Ile Tyr Gln
 100 105 110
 Lys Ile Asp Glu Ser Lys Leu Lys Pro Tyr Thr Arg Tyr Leu Val Arg
 115 120 125

<213> *Bacillus thuringiensis*

<400> 27

Met	Phe	Ile	Ser	Asn	Ile	Lys	Asn	Thr	Leu	Lys	Ile	Glu	Thr	Thr	Asp	1	5	10	15
Tyr	Glu	Ile	Asp	Gln	Ala	Ala	Ile	Ser	Ile	Glu	Cys	Met	Ser	Asn	Glu	20	25	30	
His	Ser	Ser	Lys	Glu	Glu	Met	Met	Leu	Trp	Asp	Glu	Val	Lys	Gln	Ala	35	40	45	
Lys	Gln	Leu	Ser	Trp	Ser	Arg	Asn	Leu	Leu	Tyr	Asn	Gly	Asp	Phe	Glu	50	55	60	
Asp	Val	Ser	Asn	Gly	Trp	Lys	Thr	Ser	Asn	Thr	Ile	Glu	Ile	Arg	Glu	65	70	75	80
Asn	Ser	Pro	Val	Phe	Lys	Gly	His	Tyr	Leu	His	Met	Phe	Gly	Ala	Arg	85	90	95	
Asp	Ile	Asp	Gly	Thr	Leu	Phe	Pro	Thr	Tyr	Ile	Tyr	Gln	Lys	Ile	Glu	100	105	110	
Glu	Ser	Lys	Leu	Lys	Pro	Tyr	Thr	Arg	Tyr	Arg	Val	Arg	Gly	Phe	Val	115	120	125	
Gly	Ser	Ser	Lys	Asp	Leu	Lys	Leu	Met	Val	Thr	Arg	Tyr	Gly	Lys	Glu	130	135	140	
Ile	Asp	Ala	Met	Met	Asn	Val	Pro	Asn	Asp	Leu	Ala	Tyr	Met	Gln	Pro	145	150	155	160
Thr	Pro	Ser	Cys	Gly	Asp	Ser	Arg	Cys	Glu	Ser	Ser	Ser	Arg	Tyr	Val	165	170	175	
Ser	Gln	Gly	Tyr	Pro	Thr	Pro	Val	Thr	Asp	Gly	Tyr	Ala	Ser	Gly	Arg	180	185	190	
Tyr	Ala	Cys	Gln	Ser	Asn	Arg	Gly	Thr	Lys	His	Val	Lys	Cys	His	Asp	195	200	205	
Arg	His	Pro	Phe	Asp	Phe	His	Ile	Asp	Thr	Gly	Glu	Leu	Asp	Thr	Asn	210	215	220	
Thr	Asn	Val	Gly	Ile	Asp	Val	Leu	Phe	Lys	Ile	Glu	Gly	Pro	Leu	Thr	225	230	235	240
Tyr	Ala	Thr	Leu	Gly	Asn	Leu	Glu	Val	Ile	Glu	Glu	Gly	Pro	Leu	Thr	245	250	255	
Gly	Glu	Ala	Leu	Thr	His	Val	Lys	Gln	Lys	Glu	Lys	Lys	Trp	Lys	Gln	260	265	270	
His	Met	Glu	Lys	Lys	Arg	Trp	Glu	Thr	Gln	Gln	Ala	Tyr	Asp	Pro	Ala	275	280	285	
Lys	Gln	Ala	Val	Asp	Ala	Leu	Phe	Thr	Asn	Glu	Gln	Glu	Leu	His	Tyr	290	295	300	
His	Ile	Thr	Leu	Asp	His	Ile	Gln	Asn	Ala	Asp	Arg	Leu	Ile	Gln	Ala	305	310	315	320
Ile	Pro	Tyr	Val	Tyr	His	Ala	Trp	Leu	Pro	Asp	Ala	Pro	Gly	Met	Asn	325	330	335	
Tyr	Asp	Gly	Tyr	Gln	Gly	Leu	Asn	Ala	Arg	Ile	Met	Gln	Ala	Tyr	Asn	340	345	350	
Leu	Tyr	Asp	Ala	Arg	Asn	Val	Ile	Thr	Asn	Gly	Asp	Phe	Thr	Gln	Gly	355	360	365	
Leu	Thr	Gly	Trp	His	Ala	Ala	Gly	Lys	Ala	Met	Val	Gln	Gln	Met	Asp	370	375	380	
Gly	Ala	Ser	Val	Leu	Val	Leu	Ser	Asn	Trp	Ser	Ala	Gly	Val	Ser	Gln	385	390	395	400
Asn	Leu	His	Val	Gln	Glu	His	His	Gly	Tyr	Met	Leu	Arg	Val	Ile	Ala	405	410	415	
Lys	Lys	Glu	Gly	Pro	Gly	Lys	Gly	Tyr	Val	Thr	Met	Met	Asp	Cys	Asn	420	425	430	

Gly	Asn	Arg	Glu	Thr	Leu	Lys	Phe	Thr	Ser	Cys	Glu	Glu	Gly	Tyr	Met
		435					440					445			
Thr	Lys	Thr	Val	Glu	Val	Phe	Pro	Glu	Ser	Asp	Arg	Val	Arg	Ile	Glu
	450					455					460				
Ile	Gly	Glu	Thr	Glu	Gly	Thr	Phe	Tyr	Val	Asp	Ser	Ile	Glu	Leu	Leu
465					470					475					480
Cys	Met	Gln	Gly	Tyr	Ala	Ser	Asn	Asn	Asn	Pro	His	Thr	Gly	Asn	Met
				485					490					495	
Tyr	Gly	Gln	Ser	Tyr	Asn	Gly	Asn	Tyr	Asn	Gln	Asn	Thr	Ser	Asp	Val
			500					505					510		
Tyr	His	Gln	Gly	Tyr	Thr	Asn	Asn	Tyr	Asn	Gln	Asn	Ser	Ser	Asn	Met
		515					520					525			
Tyr	Asn	Gln	Asn	Tyr	Thr	His	Asn	Asp	Asp	Leu	His	Ser	Gly	Cys	Thr
		530				535					540				
Cys	Asn	Gln	Gly	His	Asn	Ser	Gly	Cys	Thr	Cys	Ser	Gln	Gly		
545					550					555					

<210> 28

<211> 558

<212> PRT

<213> Bacillus thuringiensis

<400> 28

Met	Phe	Thr	Asn	Gly	Thr	Lys	Asn	Thr	Leu	Lys	Ile	Glu	Thr	Thr	Asp
1				5					10					15	
Tyr	Glu	Ile	Asp	Gln	Ala	Ala	Ile	Ser	Ile	Glu	Cys	Met	Ser	Asp	Glu
			20					25					30		
His	Ser	Pro	Lys	Glu	Lys	Met	Met	Leu	Trp	Asp	Glu	Val	Lys	Arg	Ala
		35				40						45			
Lys	Leu	Leu	Ser	Gln	Ser	Arg	Asn	Leu	Leu	Gln	Asn	Gly	Asp	Phe	Gly
	50					55				60					
Asp	Phe	Tyr	Gly	Asn	Asp	Trp	Lys	Phe	Gly	Asn	Asn	Ile	Ile	Ile	Gly
65					70					75					80
Ser	Asn	Asn	Ser	Ile	Phe	Lys	Gly	Asn	Phe	Leu	Gln	Met	Ser	Gly	Ala
				85					90					95	
Arg	Asp	Ile	Tyr	Gly	Thr	Ile	Phe	Pro	Thr	Tyr	Ile	Tyr	Gln	Lys	Ile
			100					105					110		
Asp	Glu	Ser	Lys	Leu	Lys	Pro	Tyr	Thr	Arg	Tyr	Arg	Val	Arg	Gly	Phe
		115					120					125			
Val	Gly	Ser	Ser	Lys	Asp	Leu	Arg	Leu	Met	Val	Thr	Arg	Tyr	Gly	Lys
	130					135					140				
Glu	Ile	Asp	Ala	Met	Met	Asn	Val	Pro	Asn	Asp	Leu	Ala	Tyr	Met	Gln
145					150					155					160
Pro	Asn	Pro	Ser	Cys	Gly	Asp	Ser	Arg	Cys	Glu	Ser	Ser	Ser	Gln	Tyr
				165					170					175	
Val	Ser	Gln	Gly	Tyr	Pro	Thr	Pro	Thr	Asp	Gly	Tyr	Ala	Pro	Asp	Arg
			180					185					190		
Tyr	Ala	Cys	Pro	Ser	Ser	Ser	Asp	Lys	Lys	His	Val	Met	Cys	His	Asp
		195					200					205			
Arg	His	Pro	Phe	Asp	Phe	His	Ile	Asp	Thr	Gly	Glu	Leu	Asp	Thr	Asn
		210				215					220				
Thr	Asn	Val	Gly	Ile	Asp	Val	Leu	Phe	Lys	Ile	Ser	Asn	Pro	Asp	Gly
225					230					235					240
Tyr	Ala	Thr	Leu	Gly	Asn	Leu	Glu	Val	Ile	Glu	Glu	Gly	Pro	Leu	Thr
				245					250					255	
Gly	Glu	Ala	Leu	Thr	His	Val	Lys	Gln	Lys	Glu	Lys	Lys	Trp	Lys	Gln

